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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                      Minimum DB seq length: 0
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                                                                                                                                                                                                                                                                                           Database
                                                                                                                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence:
SPTREMBL 25:*

1: sp_archea:*
2: sp_bacteria:*
3: sp_fungi:*
4: sp_human:*
5: sp_invertebrate:*
6: sp_mammal:*
7: sp_mbc:*
10: sp_blage:*
11: sp_rodent:*
11: sp_rodent:*
12: sp_virus:*
13: sp_virus:*
14: sp_unclassified:*
15: sp_bacteriap:*
17: sp_archeap:*
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Gapop 10.0 , Gapext 0.5
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117
1 QRVEILEGRTECVLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1017041 segs, 315518202 residues
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

16	15	14	13	12	11	10	9	80	7	0	ហ	4	ω	N	_	Result No.
47	47	47	48	48	48	49	49	52	56	104	104	104	104	107	109	Score
40.2	40.2	40.2	41.0	41.0	41.0	41.9	41.9	44.4	47.9	88.9	88.9	88.9	88.9	91.5	93.2	Query Match I
640	539	539	1028	1028	428	1028	826	1028	176	418	387	229	229	509	316	Length
10	10	10	11	11	10	11	N	4	17	σ	თ	σ	თ	თ	11	DB
Q9ASX3	Q7XDN4	Q94LS6	Q8C6X1	Q9JMB8	Q9LPY7	P97528	Q9AKX8	Q9UQ52	Q8TYF5	Q95N14	Q95N13	Q27950	Q28206	6ZAW6Ö	035545	ID
	Q7xdn4 oryza sativ	Q94ls6 oryza sativ	Q8c6x1 mus musculu	Q9jmb8 mus musculu	Q9lpy7 arabidopsis	P97528 rattus norv	Q9akx8 legionella	Q9uq52 homo sapien	Q8tyf5 methanopyru	Q95n14 ovis aries	Q95nl3 ovis aries	Q27950 bos indicus	Q28206 bos taurus	Q9myz9 sus scrofa	O35545 rattus norv	Description

	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17
	44	44	44	44	•	44.5	44.5	45	45	45	45	45	45	45	45	45	45	45	45.5	46	46	46	46	46	46	47	47	47	47
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•	O91632 cherry gree	drosophila	Q8z0g3 anabaena sp	Q8xfl2 salmonella	Q9zwi3 cucurbita m	O96680 drosophila	Q9vml1 drosophila	O57161 spinach lat	Q7yz47 cryptospori	Q8a0c0 bacteroides		Q95r48 drosophila	Q9u7p5 eufolliculi	Q94ct7 oryza sativ	Q9i2j3 pseudomonas	Q30747 macaca neme	Q30567 macaca fasc		Q8tf83 agrocybe ae	Q8ta82 caenorhabdi	Q82857 jembrana di		Q9cxf5 mus musculu	Q.	-		Q7ul12 rhodopirell		

ALIGNMENTS

Query	SOFT	322		R R R R	R R R	RRA	R R R R R R R R R R R R R R R R R R R	8888	BALLA	RESULT 035545 ID 0
Query Match Best Local Similarity	SIGNAL CHAIN SEQUENCE	SMART; SMO0060; FN3; 1. PROSITE; PS01352; HEMATOPO_REC_L_F1;	<pre>InterPro; IPR003961; FN_III. InterPro; IPR008957; FN_III-like. InterPro; IPR003528; Hemtopoptn_L_F1. Dfam. BB0001. fp.</pre>	GO; GO:0016020; GO; GO:0004896; GO; GO:0004872; InterPro; IPR002	Biochim. Biophys. Acta 1403:169-178(1998) EMBL; D83509; BAA22373.1; - HSSP; P19235; 1EBA.	Miyatake K., Nakano Y.; "The intron 5-inserted form of rat expressed as a membrane-bound form.	[1] - SEQUENCE FROM N.A. MEDLINE=98296111; PubMed=9630610; Yamaji R., Murakami C., Takenoshita	Rattus norvegicus (Rat). Eukaryota; Metazoa; Chor Mammalia; Eutheria; Rode NCBI TaxID=10116;	03545; 01-JAN-1998 (TrEMBLrel. 05, Created) 01-JAN-1998 (TrEMBLrel. 05, Last sequence update) 01-OCT-2003 (TrEMBLrel. 25, Last annotation update) Intron 5-inserted form of erythropoietin receptor precursor	LT 1 45 035545
ilarity	25 316 AA;	060; FN3; 01352; HE	IPR003961; FN_II IPR008957; FN_II IPR003528; Hemto		ophys. Ac 9; BAA223 5; 1EBA.	, Nakano 15-insert 18 a membr	OM N.A. 196111; Pu Murakami	regicus (R Metazoa; jutheria; il0116;	(TrEMBLrel. (TrEMBLrel. (TrEMBLrel.	PRELIMINARY;
93.2%; 91.3%;	24 316 34220 MW;	1. MATOPO_RE	FN_III. FN_III-1 Hemtopop	<pre>C:membrane; IEA. F:hematopoietin/interferon-class F:receptor activity; IEA. 996; CRIA.</pre>	ta 1403:1 73.1;	Y.; ed form o ane-bound	ıbMed=9630 C., Taken	Rat). Chordata; Rodentia;	el. 05, C el. 05, L el. 25, L	ARY;
Score 109; Pred. No.	24	C_L_F1; 1	ike. htn_L_F1.	EA. in/interf tivity;	.69-178 (19	f rat ery	0610; loshita M.,	Craniata Sciurogi	Created) Last sequence update) Last annotation update) Ythropoietin receptor p	PRT;
); DB 11; . 6.8e-10;	TENTIAL. TENTIAL. 05C44BF8516C180B			rferon-cla IEA.	998).	erythropoietin	., Tsuyama	ı; Verteb nathi; Mu	ced) sequence update) annotation updat ppoietin receptor	316 AA.
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Q28206;
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01-NOV-1996
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Q9MYZ9;
01-OCT-2000
01-OCT-2000
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GG; GO:0004896; F:hematopoietin/interferon-class (D200-domain.
GG; GO:0004872; F:receptor activity; IEA.
GG; GO:0004872; F:receptor activity; IEA.
GG; GO:0006118; P:electron transport; IEA.
InterPro; IPR002996; CRIA.
InterPro; IPR003961; FN III.
InterPro; IPR003961; FN III-like.
InterPro; IPR003957; FN III-like.
InterPro; IPR003528; Hemtopoptn L.F1.
InterPro; IPR003528; Hemtopoptn L.F1.
InterPro; IPR003528; Hemtopoptn L.F1.
InterPro; IPR00357; Oxidored_molyb.
BMART; SM00060; FN3; 1.
   TISSUE=Bone marrow;
Suliman H.B., Feldman
Submitted (JUN-1996) t
                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                        EPOR.
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SEQUENCE
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PROSITE; PS00559; MOLYBDOPTERIN EUK; 1.
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Christenson R.K., Vallet J.
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                          Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                                                                      Erythropoietin
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[1]
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                                                                                                SEQUENCE FROM N.A.
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(TrEMBLrel. 01, Last sequence up
(TrEMBLrel. 25, Last annotation
in receptor (Fragment).
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   Majiwa P.A.O., Logan-Henfrey EMBL/GenBank/DDBJ databases.
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Pred. No. 2.
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.5e-09;
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Sus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Expression
                                                                                                                                                                                                                              Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Best Local
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Best Local
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                                                                             Receptor.
NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                             01-NOV-1996
01-NOV-1996
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS01352; I
Receptor.
NON_TER 1
NON_TER 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR003961; FN_III.
InterPro; IPR008957; FN_III-like.
InterPro; IPR003528; Hemtopoptn_L
Pfam; PF00041; fn3; 1.
SMART; SM00060; FN3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
                                                                                                                                                                    GO; GO:0016020; C:membrane; IEA.
GO; GO:0004896; F:hematopoietin/interferon-class
GO; GO:0004872; F:receptor activity; IEA.
InterPro; IPR003996; CR1A.
InterPro; IPR003961; FN_III.
                                                                                                                                                                                                                      Suliman H.B., Feldman B.F., Submitted (JUN-1996) to the EMBL; U61398; AAB03870.1; -. HSSP; P19235; 1EBA.
                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Ce
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                    Erythropoietin receptor
                                                                                                                                                                                                                                                                                                                                                                                            Q27950;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GO; GO:0016020; C:membrane; IEA.
GO; GO:0004896; F:hematopoletin/interferon-class (D200-domain.
GO; GO:0004872; F:receptor activity; IEA.
InterPro; IPR002996; CR1A.
                                                                                                                     PROSITE;
                                                                                                                                SMART;
                                                                                                                                        Pfam; PF00041; fn3;
                                                                                                                                                                                                                                                             TISSUE=Bone marrow;
                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                  Bos indicus (Zebu).
                                                                                                                                                                                                                                                                                                                                                                                                      Q27950
                                                                                                                                                                                                                                                                                         NCBI_TaxID=9915;
                                                                                                                                                   InterPro;
                                                                                                                                                            interPro;
                                                                                                                                                                                                                                                                                                                                                                                                                         4
                                                                                                                                                                                                                                                                                                                                                                                                                                                       136
 136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              U61399; AAB03871.1;
P19235; 1EBA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20;
                                                                                                                                SM00060;
                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                      PS01352;
ORVEILDGRTECLLSNIRGGTRY 158
            QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                       QRVEILDGRTECLLSNLRGGTRY 158
                                                                                                                                                   IPR003528;
                                                                                                                                                             IPR008957;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      229
229 AA;
                                                                              229
229 I
                                                                                                                                                                                                                                                                                                                                                             (TrEMBLrel.
                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                              ΑA;
                                                                                                                                FN3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HEMATOPO_REC_L_F1;
                                                                             1
229
25196 MW;
                                                                                                                   HEMATOPO_REC_L_F1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    229
25170 MW;
                                                                                                                                                                                                                                                                                                              Chordata; Craniata; Vertebrat
Cetartiodactyla; Ruminantia;
                                                                                                                                                                                                                                                                                                                       Chordata;
                                               88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88.9%;
                                                                                                                                                Hemtopoptn_L_F1
                                                                                                                                                           FN_III.
FN_III-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hemtopoptn_L_F1.
                                                                                                                                                                                                                                                                                                                                                    01, Created
01, Last se
25, Last ar
(Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Score 104; DB; Pred. No. 3.3e. 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation updat
                                      2;
                                      Score 104; DB Pred. No. 3.3e 2; Mismatches
                                                                                                                                                                                                                                          Majiwa P.A.O., Logan-Henfrey EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                 Created)
                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                              F6E01C4AB07893E8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EDFAA6F110D992E8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                       229
                                                                                                                                                                                                                                                                                                              Vertebrata; Euteleostomi;
minantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6;
.3e-09;
                                                 .3e-09
                                                          DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                       A
                                                                                                                                                                                                                                                                                                                                                               update)
                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                   (D200-domain.
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                                       Indels
                                                          229;
                                                                                                                                                                                                                                                   L.L.;
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                                       Gaps
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RESULT Q95N13

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RESULT 6
Q95N14
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Best Local S
Matches 20
                                                                                                                                                                                                     SEQUENCE FROM N.A.

David B., Lim G.B., Moritz K.M., Koukoulas I., Winto David B., Lim G.B., Moritz K.M., Koukoulas I., Winto David B., Lim G.B., Moritz K.M., Koukoulas I., Winto David B., Lim G.B., Moritz K.M., Koukoulas I., Winto David B., Moritz K.M., Koukoulas I., Winto David B., Winto David B., Moritz K.M., Koukoulas I., Winto David B., Winto 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q95N13
Q95N13;
01-DEC-2001
01-DEC-2001
01-OCT-2003
Receptor.
NON_TER
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q95N14 PRELIMINARY; PRT; 418 AA. Q95N14; Q95N14; O1-DEC-2001 (TrEMBLrel. 19, Created) O1-DEC-2001 (TrEMBLrel. 19, Last sequence update) O1-OCT-2003 (TrEMBLrel. 25, Last annotation updat Erythropoietin receptor (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ovis aries (Sheep).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
Bovidae; Caprinae; Ovis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the ovine fetus.";
Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AY029232; AAK38737.1; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0004896; F:hematopoietin/interferon-class (D200-domain.
GO; GO:0004872; F:receptor activity; IEA.
                                                                                                                         InterPro; IPR003528; H
Pfam; PF00041; fn3; 1.
SMART; SM00060; FN3; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00041; fn3; 1. SMART; SM00060; FN3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR003961; FN III.
InterPro; IPR008957; FN III-like.
InterPro; IPR003528; Hemtopoptn L F1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. David B., Lim G.B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9940;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ovis aries (Sheep)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Erythropoietin receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bovidae; Caprinae; Ovis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9940;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Quantitation of the
                                                                                               PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ORVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                               PS01352;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORVEILDGRTECLLSNIRGGTRY 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PS01352; HEMATOPO_REC_L_F1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1
387
387 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (TrEMBLrel. 19, Created)
(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
in receptor (Fragment).
      1
418
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                                                                                         HEMATOPO_REC_L_F1; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1
387
      1
418
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e mRNA levels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     42039 MW;
                                                                                                                                                                            Hemtopoptn_L_F1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 104; DB
Pred. No. 5.9e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EpoR
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                                                                                                                                                                                                                                                                                                                                                                                                 databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wintour M.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 387;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CRC64;
                                                                                                                                                                                                                                                                                                           (D200-domain.
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RESULT
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AC Q5
AC Q5
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Q8TYF5;
Q1-JUN-2002
01-JUN-2002
01-OCT-2003
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01-MAY-2000
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01-OCT-2003
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STRAIN=AV19 / DSM 6324 / JCM 9639;

MEDLINE=21927647; PubMed=11930014;

Slesarev A.I., Mezhevaya K.V., Makarova K.S., Polushin Shcherbinina O.V., Shakhova V.V., Belova G.I., Aravind Natale D.A., Rogozin I.B., Tatusov R.L., Wolf Y.I., Ste Malykh A.G., Koonin E.V., Kozyavkin S.A., While Complete genome of hyperthermophile Methanopyrus k and monophyly of archaeal methanogens.";

Proc. Natl. Acad. Sci. U.S.A. 99:4644-4649(2002).
EMBL; AB003592; BAAB2612.1; -. HSSP; P20241; 1CFB. Genew; HGNC:2176; CNTIN6. GO; GO:0007155; P:cell adhesion; TAS.
                                                                                                                         SEQUENCE FROM N.A.

MEDILINE-98146168; PubMed-9486763;

Kamei Y., Tsutsumi O., Taketani Y., Watanak

"cDNA cloning and chromosomal localization

molecule, NB-3 in human.";

J. Neurosci. Res. 51:275-283 (1998).
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EMBL;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01.MAY-2000 (TrEMBLrel. 13, Created)
01.MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CCT-2003 (TrEMBLrel. 25, Last annotation updat
Neural adhesion molecule NB-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9UQ52
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NCBI_TaxID=2320;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human)
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InterPro; IPR009003; Cys_Ser_trypsin.
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Primates;
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Pred. No.
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Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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P97528;
01-MAY-1997
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InterPro; IPR003
InterPro; IPR003
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Q9AKX8;
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Pfam; PF00047; f9; 6.
SMART; SM00060; FN3; 3.
SMART; SM00060; FN3; 3.
SMART; SM00408; IGC2; 5.
PROSITE; PS00211; ABC TRANSPORTER 1; 1.
PROSITE; PS0835; IG_LIKE; 6.
                                                                                                                                        NB-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lueneberg E., Mayer B., Daryab N., Kooistra O., Zaehringer U., Rohde M., Swanson J., Frosch M.; "Chromosomal insertion and excision of a 30 kb instable genetic element is responsible for phase variation of lipopolysaccharide other virulence determinants in Legionella pneumophila."; Mol. Microbiol. 39:1259-1271(2001).
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InterPro; IPR003961; FN III-
InterPro; IPR0039957; FN III-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig_c2.
                              NCBI_TaxID=10116;
                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical protein.
SEQUENCE 826 AA; 91581 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=serogroup 1;
MEDLINE=21150471; PubMed=11251842;
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Legionellaceae; Legionella.
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    SEQUENCE FROM N.A
                                                                                                            Rattus norvegicus (Rat)
                                                                                                                                                             01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypothetical protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=446;
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Local Similarity 42.9%;
nes 9; Conservative
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(TrEMBLrel. 17, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
                                                                                                                                                        (TrEMBLrel. 03, Created)
(TrEMBLrel. 03, Last seq
(TrEMBLrel. 25, Last ann
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                                                                                                                                                                                                                                                   PRELIMINARY;
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                                                                     Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             44.48;
71.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gammaproteobacteria; Legionellales;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 49;
Pred. No.
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                                                                   Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rat
                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
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Mismatches
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RESULT 11
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RT from rat brain.";

RI enterpro; Ispanding casette (ABC) transporter acti...; IEA.

RGO; GO:0004009; F:ATP-binding casette (ABC) transporter acti...; IEA.

RGO; GO:0006810; P:transport; IEA.

RI interpro; IPR00343; ABC transporter.

RI Interpro; IPR00396; FN III-like.

RR Interpro; IPR00396; FN III-like.

RR Interpro; IPR00359; FN III-like.

RR Interpro; IPR00359; FN III-like.

RR Interpro; IPR00359; IG_C2.

RR SMART; SM00060; FN3; 4.

RR SMART; SM00408; IGC2; 5.

RR SMART; SM00408; IGC2; 5.

RR PROSITE; PS0031; IG_LIKE; 6.

RR FROSITE; PS0031; IG_LIKE; 6.
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Best Local S
Matches 10
                                                                               Best
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01-OCT-2000
01-OCT-2000
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                                                                                                                                                                                                              Shinn P., Brooks S., Buehler E., Chao Q., Johnson-Hopson C., Kh Kim C., Altafi H., Bei Q., Chin C., Chiou J., Choi E., Conn L., Conway A., Gonzales A., Hansen N., Howing B., Koo T., Lam B., Lienz C., Li J., Liu A., Liu K., Liu S., Mukharsky N., Nguyen M. Palm C., Pham P., Sakano H., Schwartz J., Southwick A., Thaveri Toriumi M., Vaysberg M., Yu G., Federspiel N.A., Theologis A., Ecker J.R.;
                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;

eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                SEQUENCE
                                                                                                                                                Submitted (OCT-2000) to the EMBL/GenBank/DDBJ databases. EMBL; AC011661; AAF16632.1; -.
                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9LPY7
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                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=3702;
                                                                                                                                                                                                "Genomic sequence for Arabidopsis thaliana BAC T23J18 from
                                                                                Local
 132
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                                                                               Similarity
KRPQLLENTTLCILSITRGSRR
                              QRVEILEGRIECVLSNLRGRIR 22
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                                                                                                                                428 AA;
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                                                                                                                             48982 MW;
                                                                               41.0%;
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71.4%;
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Last annotation update)
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Pred. No.
                                                                               Pred.
                                                                                                Score 48;
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                                                                                                                                  603F3FF2725622F9 CRC64;
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32;
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., Thaveri A.,
                                                                                                                                                                                                  chromosome
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Best Local S
Matches 9
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Q8C6X1;
01-MAR-2003
01-MAR-2003
01-OCT-2003
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Q9JMB8;
01-OCT-2000
SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUB=Head;
MEDLINE=22354683; PubMed=12466851;
The FANTOM Consortium,
the RIKEN Genome Exploration Resea
                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                         Pfam; PF00041; fn3; 4.
Pfam; PF00047; ig; 6.
SMART; SM00060; FN3; 4.
SMART; SM00408; IGC2; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MGD; MGI:1858223; Cntn6.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005524; F:ATP binding; IEA.
GO; GO:0004009; F:ATP-binding cassette
GO; GO:0006810; P:transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Expression and regulation of a gene encoding molecule NB-3 of the contactin/F3 subgroup in Gene 245:253-266(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=129SVJ; TISSUE=Brain;
MEDLINE=20183687; PubMed=10717476;
Lee S., Takeda Y., Kawano H., Hosoya H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS00211; ABC_TRANSPORTER_1; PS50835; IG_LIKE; 6. PROSITE; PS00290; IG_MHC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AB032602; BAA92367.1; HSSP; P20241; 1CFB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Neural recognition molecule
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                                                                                                    Mus musculus
                                                                                                                 CNTN6.
                                                                                                                         Contactin 6.
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                                                                                                                                                                                                                                                                                                                                              Immunoglobulin domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ; IPR003439; ABC transport

; IPR003961; FN III-

; IPR008957; FN III-11ke.

; IPR007110; Ig-11ke.

; IPR003598; Ig_c2.

; IPR003006; Ig_MHC.
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(TIEMBLIE) 15, Last sequence update)
(TIEMBLIEL 25, Last annotation update)
phition molecule NB-3.
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                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                 PRELIMINARY;
                                                                                                    (Mouse)
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,
 Exploration Research Group Phase I & II Team;
                                                                             Chordata;
Rodentia;
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Pred. No. 47;
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                                                                             Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
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GO; GO:0005524; F:ATP binding; IEA.
GO; GO:0004009; F:ATP-binding cassette
GO; GO:0006810; P:transport; IEA.
InterPro; IPR003439; ABC transporter.
InterPro; IPR003957; FN_III.
InterPro; IPR003957; FN_III.1ike.
InterPro; IPR003599; Ig.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q94LS6;
01-DEC-2001
01-DEC-2001
                                                                                                                                                                                                                    STRAIN=cv. Nipponbare;
Buell C.R., Yuan Q., Ouyang S., Moffat K.S., Hill J.N., Gansber;
Brenner M., Burgess S., Hance M., Shvartsbeyn M., Tsitrin T.,
Riggs F., Hsiao J., Zismann V., Blunt S., Pai G., Vanaken S.E.,
Utterback T.R., Feldblyum T.V., Quackenbush J., Salzberg S.L.,
White O., Fraser C.M.;
"Oryza sativa chromosome 10 BAC OSJNBb0011A08 genomic sequence.
Submitted (MAY-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                               Hypothetical
SEQUENCE 5:
                                                                                                                                                                                                       Gramene; Q94LS6; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hypothetical protein. Oryza sativa (Rice).
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01-OCT-2003 (TrEMBLrel.
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Nature 420:563-573(2002)
                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR003598;
InterPro; IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Analysis of the mouse transcriptome
                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=4530;
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Last annotation update)
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RESULT 15

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O7XDN4
ID O7XDN4
PRELIMINARY; PRT; 539 AA.

AC Q7XSN4;
DT O1-OCT-2003 (TrEMBLrel. 25, Last sequence update)
DT O1-OCT-2003 (TrEMBLrel. 25, Last sequence update)
DT O1-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DT O1-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Hypothetical protein.
GN OSUNBB0011A08.11.
OS Oryza sativa ()aponica cultivar-group).
OC Enkaryora; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
OC Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
OX NCB1_TaxID=3947;
RN [1]
RP SEQUENCE FROM N.A.
The Rice Chromosome 10 Sequencing Consortium;
RA The Rice Chromosome 10 Sequencing Consortium;
RI "In-depth view of structure, activity, and evolution of rice chromosome 10.";
RL Science 300:1566-1569(2003).
RN [2]
RP SEQUENCE FROM N.A.
SCOTRAIN=cv. Nipponbare;
RA Buell C.R., Wing R.A., McCombie W.R., Messing J., Yuan Q.;
SUMML; Asg017102; AAP54107.1; -.
KW Hypothetical protein.
SQ SEQUENCE 539 AA; 60201 MW; 8EBED40239310BC2 CRC64;
Search completed: May Job time : 40 secs
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STRAIN-cv. Nipponbare;
STRAIN-cv. Nipponbare;
Buell C.R., Wing R.A., McCombie W.R., Messing J., Yuan Q.;
Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AE017102; AAP54107.1; -.
Hypothetical protein.
SEQUENCE 539 AA; 60201 MW; 8EBED40239310BC2 CRC64;
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RESULT 1

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ID EPORR HUMAN
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DT 01-NOV-1990
RX EXPLINE=903
RA JONES S.S.,
RT "HUMAN erythry
RY MEDLINE=903
RA JONES S.S.,
RT "HUMAN EY-THY
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RY MEDLINE=903
RA MEDLINE=913
RA MEDLINE=913
RA HANKINE W.D
RT "Cloning of RL Blood 78:25
RN [4]
RP SEQUENCE OF RC TISSUE=Plac
RX MEDLINE=923
RA MAOUCHE L.,
RA MEDLINE=923
RA MAOUCHE L.,
RA Chretien S.
RT "Cloning of RL Blood 78:25
RN [4]
RP SEQUENCE OF REDLINE=921
RA MAOUCHE L.,
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RA MAOUCHE L.,
RA MAOUCHE L.,
RA CHRETIEN S.
RT "Cloning of RL Blood 78:25
RN [4]
RP SEQUENCE OF REDLINE=921
RA MAOUCHE L.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIINE=91372359; PubMed=1654273;
Ehrenman K., St John T.;
"The erythropoletin receptor gene: cloning and
multiple transcripts in an erythroid cell line
Exp. Hematol. 19:973-977(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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01-NOV-1990 (Rel. 16, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation updat
Erythropoietin receptor precursor (EPO-R).
                                                                                                                                                                                                                                                                                                           Chretien S.;
"Cloning of the gene encoding Blood 78:2557-2563(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Cloning of the human erythropoietin receptor gene."; Blood 78:2548-2556(1991).
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MEDLINE=92399733; PubMed=1668606;
NOGUCHI C.T., Bae K.S., Chin K., Wada Y., Schechter A.N.,
Hankins W.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=90304340; PubMed=2163696; Jones S.S., D'Andrea A.D., Halnes L.L., Williams erythropoletin receptor: cloning, characterization.";
                                                                                                                                                                                                                                                                 SEQUENCE OF 1-17 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=92399734; PubMed=1668607;
Maouche L., Tournamille C., Hattab C., Boffa G., Cartron J.-P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Placenta;
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  "APS, an adaptor homology-2 (SH2)
                                                                         Wakioka T.,
                                                                                             PHOSPHORYLATION, I
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                                              foshimura A.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STANDARD;
  protein containing Pleckstrin homology (PH) and domains inhibits the JAK-STAT pathway in
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GLAR_ECOLI
XYNB_THENE
PROA_CAUCR
Y997_ARCFU
YHX7_YEAST
PAX3_HUWAN
PAX3_MOUSE
PAX7_HUWAN
HS7F_CAEEL
DPO1_THEFI
PT1A_ECOLI
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                                                                       Inoue A., Komiya S.,
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Q60046
O29256
O29267
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P3860
P24610
P23759
P11141
O52225
P32670
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biologic

Of.

Result No.

Score

Query Match

Length

BB

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SUMMARIES

EPOR_HUMAN EPOR_MOUSE EPOR_RAT GIDB_BIFLO KRM1_MOUSE KRM1_HUMAN

PYRB_COREF PYRB_CORGL TSP3 HUMAN YS59_ENTFA RS28_KLULA RS28_KLULA RS28_KLULA RS28_KLUMA C128_MYCTU AMS2_SCHPO YL54_CAEEL DB10_NICSY YR33_MYCTU AXO1_CHICK KRMI_RAT FPG_VIBPA MURG_CORGL TSP3_MOUSE YA29_TREPA PTED_HUMAN RS28_YEAST DHAS_YEAST DHA

41.9 41.0 41.0 39.3 39.3 39.3 39.3

42.5 42.5 42.4 42.4 42.4 42.4 42.5 41.5

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GenCore version (c) 1993 - 2004

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EMBL; M34986; AAA52401.1; EMBL; M60459; AAA52403.1; EMBL; M60459; AAA52393.1; EMBL; M75595; AAA52393.1; EMBL; M77544; AAA52392.1; EMBL; M77544; AAA52392.1; EMBL; M77544; AAA52392.1; EMBL; M77544; AAA52392.1; EMBL; AA3799; ZUUL-97. PDB; 1EBA; 29-UUL-97. PDB; 1EBA; 18-NOV-98. PDB; 1CM4; 11-AUG-99. PDB; 1CM4; 
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Syed R.S., Reid S.W., Li C., Cheetham J.C., Aoki K.H., Liu B.
Zhan H., Osslund T.D., Chirino A.J., Zhang J., Finer-Moore J.
Elliott S., Sitney K., Katz B.A., Matthews D.J., Wendoloski Cegrie J., Stroud R.M.;
"Efficiency of signalling through cytokine receptors depends
critically on receptor orientation.";
PDB; 1CNN; 11-AUG-99.
PDB; 1ENN; 07-JAN-00.
PDB; 1ENN; 07-JAN-00.
Genew; HGNC:3416; EPOR.
MIM; 133171; -.
GO; GO:000587; C:integral to plasma membrane;
GO; GO:0004900; F:erythropoietin receptor activ
InterPro; IPR002996; CRIA.
InterPro; IPR008957; FN_III-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Crystallographic evidence for preformed receptor before ligand activation."; Science 283:987-990(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              X-RAY CRYSTALLOGRAPHY (2.7 ANGSTROMS) OF MEDLINE=99023198; PubMed=9808045; Livnah O., Johnson D.L., Stura E.A., Farr You Y., Liu K.D., Goldsmith M.A., He W., Jolliffe L.K., Wilson I.A.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               X-RAY CRYSTALLOGRAPHY (2.40 ANGSTROMS) OF 34-246.
MEDLINE=99141272; PubMed=9974392;
Livnah O., Stura E.A., Middleton S.A., Johnson D.L.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entities requires a license agreement (Se or send an email to license@isb-sib.ch).
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Leukemia 13:760-767(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "An
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induced erythroblast proliferation and differentiation. Upon EPO
stimulation, EPOR dimerizes triggering the JAK2/STAT5 signaling
cascade. Isoform 2 acts as a dominant-negative receptor of EPOR-
mediated signals.
SUBURIT: Interacts with APS.
SUBURIT: Interacts with APS.
SUBCELLULAR LOCATION: Type I membrane protein.
SIMILARITY: Belongs to the type I cytokine family of receptors.
Subfamily 1.
SIMILARITY: Contains 1 fibronectin type III domain.
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. Struct. Biol. 5:993-1004(1998).
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W., Krause C.D.,
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                                                                   activity;
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EPOR MOUSE STANDARD; PRT; 50, 7...
P14753; Q63852;
01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
15-MAR-2004 (Rel. 43, Dest annotation update)
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PIRSF; PIRSF001959; EPO_receptor;
SMART; SM00060; FN3; 1.
     Eukaryota; Metazoa;
Mammalia; Eutheria;
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InterPro; IPR003528; Hemtopo
                Mus musculus (Mouse)
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                                                                                                              Similarity
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240
508
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mbrane; Glycoprotein;
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          Chordata;
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                                                                                                              Score 117;
Pred. No. 8.
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N-LINKED (GLCNAC.
                                                                                                                                                                                                                                                                                                                                                      EXTRACELLULAR (POTENTIAL).
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     Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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D'Andrea A.D., Lodish H.F., Wong G.G.; "Expression cloning of the murine erythropoietin receptor."; Cell 57:277-285(1989).
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or send a
EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Spleen focus-forming virus long terminal activation of the murine erythropoietin refriend leukemia cell line.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=92017832; PubMed=1656233;
Hino M., Tojo A., Misawa Y., Morii H., Takaku F., Shibu
"Unregulated expression of the erythropoietin receptor
insertion of spleen focus-forming virus long terminal r
interceptor of spleen focus forming virus long terminal representation of spleen focus forming virus long terminal representations.";
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STRAIN=BALB/c; TISSUE=Liver;
MEDLINE=91080149; PubMed=2175360;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "The adapter protein APS associates with the multifunctional docking sites Tyr-568 and Tyr-936 in c-Kit."; Biochem. J. 370:1033-1038(2003).
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"Characterization of murine erythropoietin
J. Mol. Biol. 216:567-575(1990).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=93180826; PubMed=8382775; Miura O., Cleveland J.L., Ihle J.N.;
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SEQUENCE FROM N.A.
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                                                                                                s SWISS-PROT entry is copyright. It is produced through a collaboratio ween the Swiss Institute of Bioinformatics and the EMBL outstation EDUTOPEN BIOINFORMATICS INSTITUTE. There are no restrictions on it by non-profit institutions as long as its content is in no waified and this statement is not removed. Usage by and for commercialities requires a license agreement (See http://www.isb-sib.ch/announce
                                                                                                                                                                                                                                                                                                                                   FUNCTION: Receptor for erythropoietin. Mediates erythropoietin-
induced erythroblast proliferation and differentiation. Upon EPC
stimulation, EPOR dimerizes triggering the JAKZ/STATS signaling
cascade. Isoform 2 acts as a dominant-negative receptor of EPOR
mediated signals.
SUBUNIT: Interacts with APS.
SUBCELLULAR LOCATION: Type I membrane protein.
SIMILARITY: Belongs to the type I cytokine family of receptors.
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  J04843;
X53081;
                                                                          an email to license@isb-sib.ch).
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  AAA37571.1;
CAA37248.1;
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Zon L.I., Orkin S.H.,
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InterPro; IPR008957; FN_III_like.
InterPro; IPR003961; FN_III.
InterPro; IPR003961; Hemtopoptn_L_F1.
Pfam; PF00041; fn3; 1.
PIRSF; PIRSF001959; EPO_receptor; 1.
SMART; SM00060; FN3; 1.
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PIR; S14081; S14081.
HSSP; P19235; 1EBA.
 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
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                                                                                                              J. Biol. Chem. -:- FUNCTION:
                                                                                                                                  cells."
                                                                                                                                           "Functional erythropoietin I characteristics. Comparison
                                                                                                                                                              Masuda S., Nagao M.,
Tabira T., Sasaki R.;
                                                                                                                                                                         MEDLINE=93266574; PubMed=7684373; Masuda S., Nagao M., Takahata K.,
                                                                                                                                                                                                                                                                     Erythropoietin receptor precursor
                                                                                                                                                                                                                                                                                                              Q07303;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                       Eukaryota;
                                                                                                                                                                                                                                                 Rattus norvegicus (Rat).
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                                     cascade. Isoform 2 acts
mediated signals.
- SUBCELLULAR LOCATION: T
- SIMILARITY: Belongs to
Subfamily 1.
                               SIMILARITY: Contains 1 fibronectin type III domain.
                                                                              FUNCTION: Receptor for erythropoietin. Mediates erythropoietin-induced erythroblast proliferation and differentiation. Upon EPO stimulation, EPOR dimerizes triggering the JAK2/STAT5 signaling cascade. Isoform 2 acts as a dominant-negative receptor of EPOR.
                                                                                                                                                                                                                                                                                                                         RAT
                                                                                                                                                                                                                   TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  M62360;
S59388;
                                                                                                                                                                                                                                                                                                                                                                            193
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                                                                                                                                                                                                                                                                                                                                                                            ORVEVLEGRIECVLSNIRGGIRY 215
                                                                                                                                                                                                                                                                                                                                                                                              QRVEILEGRIECVLSNLRGRIRY 23
                                                                                                                                                                                                                            Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transmembrane;
                                                                                                                                                                                                                                       Metazoa;
                                                                                                                                                                                                                                                                               (Rel. 31, Created)
(Rel. 31, Last sequence update)
(Rel. 43, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               250
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; AAA37572.1;
; AAA37582.1;
; AAB20029.2;
                                                                                                                       268:11208-11216(1993)
                                                                                                                                                                                                                                                                                                                         STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HEMATOPO REC L F1; 1 nbrane; Glycoprotein;
                                                                                                                                                                                                                             Chordata;
Rodentia;
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n with rec
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L; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                            Score 109;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-LINKED (GLCNAC. . .) (POTEN E -> D (IN REF. 3; AAB20029).
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BY SIMILARITY
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APS-BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             POTENTIAL.
CYTOPLASMIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EXTRACELLULAR (
                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Rattus.
                                                 I membrane
type I cyto
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                                                                                                                                           receptor
                                                                                                                                                                          Konishi Y.,
                                                                                                                                                                                                                                                                      (EPO-R)
                                                                                                                                                       of the
                                                                                                                                                                                                                                                                                                                         507
                                                 cytokine far
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of erytl
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                                                                                                                                             erythroid
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GIDB_BIFLO
Q8G6J4;
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Receptor;
This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation between Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-OCT-2003 (Rel. 42, 0
10-OCT-2003 (Rel. 42, 1
10-OCT-2003 (Rel. 42, 1
Methyltransferase gidB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                    MEDLINE=22294977; PubMed=12381787;
Schell M.A., Karmirantzou M., Snel B., Vilanova D., Berg Schell M.A., Larvier F., Bork P., Delley M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bifidobacterium longum.
Bacteria; Actinobacteria; Actinobact
Bifidobacteriaceae; Bifidobacterium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BIFLO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; D13566; BAA02761.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a
                                                                                                                                                                                                                                                               Ö
                                                                                                                                                                                                                                                                                           Pridmore R.D., Arigoni
                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=NCC 2705;
                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GIDB OR BL0646.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein B)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=216816;
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                                                                                                                                                                                                                                                               the
                                                                                                                                                                                     e genome sequence of Bifidobacterium longum refl
the human gastrointestinal tract. ",
c. Natl. Acad. Sci. U.S.A. 99:14422-14427(2002).
FUNCTION: probable S-adenosyl-L-methionine depenantly transferase specific for a sterol and/or
                                                                                                                                                  SIMILARITY: Belongs to the gidB family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A46713; A46713.
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Pro; IPR002996; CR1A.
Pro; IPR008957; FN III-like.
Pro; IPR003961; FN III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SM00060; FN3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PIRSF001959; EPO_receptor; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                         Zwahlen M.-C., Desiere F.,
R.D., Arigoni F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transmembrane;
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Last
B (EC
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BY SIMILAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Actinobacteridae; Bifidobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 annotation update)
2.1.-.-) (Glucose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REC_L_F1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         POTENTIAL.

CYTOPLASMIC (POTENTIAL).

FIBRONECTIN TYPE-III.

BY SIMILARITY.

BY SIMILARITY.

N'LINKED (GLCNAC. . . ) (P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 1.5
L; Mismatches
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -LINKED (GLCNAC. . .) (POTENTIAL)
AC79AF22D06A7312 CRC64;
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.5e-09;
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                                                                                                                                                                                            ine dependent
and/or lipid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inhibited division
                                                                                                                                                                                                                                                                                   reflects
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                              Berger B.,
                                                                                                                                                                                                                                                                                   its adaptation
                                                                                                                                                                                                substrate
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InterPro;

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RESULT 5
KRM1_MOUSE
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Q99N43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Brain, and Kidney;
MEDLINE=21167372; PubMed=11267660;
Nakamura T., Aoki S., Kitajima K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-FEB-2003 (Rel. 41, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Kremen protein 1 precursor (Kringle-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TIGREMAN, TIGRO0138; gidB; 1.
Transferase; Methyltransferase;
SEQUENCE 221 AA; 24163 MW; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AE014686; AAN24468.1; -. HAMAP; MF_00074; -; 1. InterPro; IPR003682; GidB. Pfam; PF02527; GidB; 1.
  MGD; MGI:1933988; Kremen.
GO; GO:0016021; C:integral to
InterPro; IPR000859; CUB.
                                                                     EMBL; AB059617;
HSSP; P00747; 10
                                                                                                                                                                                the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and the nose) (Dickkopf receptor).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or send an email to license@isb-sib.ch}
                                                                                                                                        or send an email
                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Molecular cloning and characterization of Kremen, a now kringle-containing transmembrane protein.", Biochim. Biophys. Acta 1518:63-72(2001).
-I-FUNCTION: Receptor for Dickkopf protein. Cooperates to block Wnt/beta-catenin signaling (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nakamura T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE SPECIFICITY: In the adult, widely expressed with high levels in heart, lung, kidney, skeletal muscle and testis. DEVELOPMENTAL STAGE: In the embryo, expression is first detected on day 9 and increases up to day 18. Lower levels are found in adult. At 9.5 dpc, expression is localised to the apical ectodermal ridge (AER) of the developing fore- and hindlimb buds, the telencephalon and the first brachial arch. At 10.5 dpc, expression is also observed in the myotome and in sensory tissues expression is also observed in the myotome and in sensory tissues.
GO:0016021; C:int
erPro; IPR000859;
                                                                                                                                                                                                                                                                                                                        such as the nasal pit and optic vesicle.
SIMILARITY: Contains 1 CUB domain.
SIMILARITY: Contains 1 Kringle domain.
SIMILARITY: Contains 1 WSC domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUBCELLULAR LOCATION: Type I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          120 QNVEIIRGRSDAVIQQVRKR 139
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                                                                                                                                          requires a license agreement (S
an email to license@isb-sib.ch).
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                                                                       1CEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STANDARD;
                                                                                             BAB40968.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DEVELOPMENTAL STAGE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        41.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 49;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 κ.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Takahashi T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          473
                                                                                                                                                             moved. Usage by and for commercial (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                          There are no restrictions ng as its content is in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matsumoto K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                        At 10.5 dpc, in sensory tissues
                                                                                                                                                                                                                                                        EMBL outstation
                                                                                                                                                                                                                                                             a collaboration
MBL outstation -
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RESULT KRAIL HUMAN HUMAN
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Best Local S
Matches 11
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PRODOM; PD000395; Kringle; 1.
SMART; SM00042; CUB; 1.
SMART; SM00130; KR; 1.
PROSITE; PS01180; CUB; 1.
PROSITE; PS00021; KRINGLE 1; 1.
PROSITE; PS00021; KRINGLE 2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HUMAN
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SEQUENCE
                                                                                                                                                                                                                                                          SEQUENCE FROM N.A. (ISOFORM 2).

Tashiro H., Yamazaki M., Watanabe K., Kumagai A., Itakura S.,

Pukuzumi Y., Fujimori Y., Komiyama M., Sugiyama T., Irie R.,

Otsuki T., Sato H., Wakamatsu A., Ishii S., Yamashita H.,

Kawai-Hio Y., Saito K., Nishikawa T., Kimura K., Yamashita H.,

Matsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda K., Wagatsuma M.,

Murakawa K., Kanehori K., Takahashi-Fujii A., Oshima A., Sugiyama A.,

Kawakami B., Suzuki Y., Sugano S., Nagahari K., Masuho Y., Nagai K.,
           MEDLINE=20057165; PubMed=10591208; Dunham I., Hunt A.R., Collins J.E., Clamp M., Smink L.J., Ainscough R., Bagguley C., Bailey J., Barlow K.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. (18 Nakamura T., Nakamura Submitted (OCT-2001) t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Kremen protein 1 precursor (Kringle-containing and the nose) (Dickkopf receptor).
KREMENI OR KREMEN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KRM1
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DOMAIN
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Pfam; PF00051; kringle; 1.
Pfam; PF01822; WSC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q96MŪ8; Q9ВY70;
28-FEB-2003 (Re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CARBOHYD
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                                                                                                                              SEQUENCE
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                                                                                                                                                                                                                human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IPR002889;
                                                                                                                                                                                   (OCT-2001) to the
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                                                                                                                        N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9UGS5; Q9UGU1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISOFORM nura T.;
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CYTOPLASMIC (POTENTIAL).

KRINGLE.

WSC.

CUB.

N-LINKED (GLCNAC. . .) (1

N-LINKED (GLCNA
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                                                                                                                                                                                project.";
EMBL/GenBank/DDBJ
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••
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 49;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KREMEN PROTEIN EXTRACELLULAR
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Bruskiewich R., Bearc
Almeida J.P., Babbage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         475
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 473;
                                      R., Beare D.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CRC64;
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              O.P.,
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RA Burrill W.D. Burton J. Carder C., Garten N.E., Connor R.
RA Clegg S.M., Cobley V.E., Cole C.G., Collier R.E., Connor R.
RA Clay D., Corby N.R., Colle C.G., Collier R.E., Connor R.
RA Conroy D., Corby N.R., Coville G.J., Cox A.V., Davis J., Dawson E.,
RA Dhami P.D., Dockree C., Dodsworth S.J., Durbin R.M., Ellington A.G.,
RA Evans K.L., Fey J.M., Flemling K., French L., Garner A.A.,
RA Hall R.E., Hall-Tamlyn G., Heathcott R.W., Ho S., Holmes S.,
RA Hall R.E., Jones M.C., Kershaw J., Kimberley A.M., King A.,
RA Hunt S.E., Jones M.C., Kershaw J., Kimberley A.M., King A.,
RA Hall R.E., Jangford C.F., Leversha M.A., Lloyd C., Lloyd D.M.,
RA Martyn I.D., Mashreghi-Mohammadi M., Matthews L.H., Mccann O.T.,
RA Martyn I.D., Mashreghi-Mohammadi M., Matthews L.H., Mccann O.T.,
RA Martyn I.D., Mashreghi-Mohammadi M., Matthews L.H., Mccann O.T.,
RA Mclay J., Mclaren S., McMurray A.A., Milne S.A., Mortimore B.J.,
RA Mclay J., Plumb R.M., Ramsay H., Ramsey Y., Rogers L., Ross M.T.,
RA Mallis S.H., Plumb R.M., Ramsay Y., Rogers L., Ross M.T.,
RA Walliams L., Williams S.A., Williamson H., Wilmer T.E., Wilming L.,
RA Williams L., Williams S.A., Williamson H., Saltson J.E., Wilming L.,
RA Williams L., Williams S.A., Williamson H., Mitsuyama S.,
RA Williams J., Ehrly D.R., Back S., Rogers J., Shimizu N.,
RA Williams J., Shibuya K., Yoshizaki Y., Aoki N., Mitsuyama S.,
RA Koe B.A., Chen F., Chu L., Crabtree J., Deschamps S., DoA., Do T.,
RA Monshima S., Kawasaki K., Sasaki T., Asakawa S., Kudoh J.,
RA Monshima S., Qian Y., Ray L., Ren Q., Shaull S., Sloan D., Song L.,
RA Minosh M., Jang F., Fu Y., Hu P., Hua A., Kenton S., Lai H., Lao H.I.,
RA Wang Q., Wang Y., Wang Z., White J., Wallingham D., Song L.,
RA Wilson R., Johnson D., Bemis G., Bentley D., Bradshaw H., Bourne S.,
RA Cordes M., Du Z., Falton L., Goela D., Graves T., Hawkins J.,
RA Wilson R., Bentley D., Shaull S., Walter T.,
RA Wilson R., Emanuel B.S., Shaihh T., Kurahashi H., Saitta S.,
RA Wilson R., Walter C., Wang Y., Walter J., Walter S.,
Ra Man M., Shang R., Yoshi
                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                       or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SIMILARITY: Contains 1 CUB domain.
-!- SIMILARITY: Contains 1 kringle domain.
-!- SIMILARITY: Contains 1 WSC domain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: Receptor for Dickkopf protein. Cooperates with Dickkopf to block Wnr/beta-catenin signaling (By similarity).
SUBCELLULAR LOCATION: Type I membrane protein (Potential).
ALTERNATIVE PRODUCTS:
Event-Alternative splicing; Named isoforms=2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IsoId=Q96MU8-1; Sequence=Displayed;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IsoId=Q96MU8-2; Sequence=VSP_003900
                                                                                                                                                                                                                                                                    an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Blakey S.E., Bridgeman A.M., Buck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    confirmation available;
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EMBL; Z95116; CAB62952.1; -.
EMBL; AL021393; CAB62959.1; -.
Genew; HGNC:17550; KREMENI.
GO; GO:0016021; C:integral to membrane; IS:
GO; GO:0005624; C:membrane fraction; TAS.
GO; GO:0007154; P:cell communication; TAS.
InterPro; IPR000859; CUB.
InterPro; IPR000859; WSC.

AB059618; BAB40969.1; -. AK056425; BAB71180.1; -.

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SOUTH THE TENT OF 
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Best Local
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TRANSMEM
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Q8FT39;
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CARBOHYD
VARSPLIC
                                                                                                                                                                            SEQUENCE FROM N.A.

STRAIN-YS-314 / AJ 12310 / DSM 44549 / JCM 11189;

MEDLINE-22723752; PubMed=12840036;

Nishio Y., Nakamura Y., Kawarabayasi Y., Usuda Y.,

Sugimoto S., Matsui K., Yamagishi A., Kikuchi H.,

Gojobori T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COREF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CONFLICT
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PROSITE; PS00021; KRINGLE_1; 1.

PROSITE; PS50070; KRINGLE_2; 1.

What signaling pathway; Signal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRINTS; PRO0018; KRINGLE.
ProDom; PD000395; Kringle;
SMART; SM00042; CUB; 1.
SMART; SM00130; KR; 1.
   This
                                                                                                                       replacements responsible efficiens.";
                                                                                                                                                                                                                                                                                                                     Bacteria; Actinobacteria; Actinobacteri
Corynebacterineae; Corynebacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                          transcarbamylase) (ATCase).
PYRB OR CE1732.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                         Corynebacterium efficiens.
                                                                                                                                                                                                                                                                                                                                                                                                           Aspartate carbamoyltransferase (EC
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                                                                                                                                                            "Comparative complete
                                                                                                                                                                                                                                                                                                    NCBI_TaxID=152794;
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                                 PATHWAY: Pyrimidine biosynthesis; second step. SIMILARITY: Belongs to the ATCase/OTCase family.
                                                                     CATALYTIC ACTIVITY: Carbamoyl phosphate + L-aspartate + N-carbamoyl-L-aspartate.
 SWISS-PROT entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COREF
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PF01822;
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475 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Rel.
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   18
                                                                                                                                      genome sequence analysis of the amino acid
ble for the thermostability of Corynebacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Created)
 copyright.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MW;
                                                                                                                                                                                                                                                                                                                      Actinobacteridae; Actinomycetales; bacteriaceae; Corynebacterium.
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KREMEN PROTEIN :

EXTRACELLULAR ()
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N-LINKED
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Pred. No.
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I -> V (IN REF. 2).
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4.1;
                                                                                                                                                                                                              Usuda Y.,
produced through a collaboration
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RESULT 8
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                                                                                             HAMAP; MF_00001; -; 1.
InterPro; IPR006130; E
InterPro; IPR006131; C
InterPro; IPR006131; C
InterPro; IPR006132; C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-2003
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InterPro; IPR006130; Asp_Orn_COtranf.
InterPro; IPR002082; Asp_carbmltransf.
InterPro; IPR006131; OTCace_O.
InterPro; IPR006131; OTCace_P.
Pfam; PF00185; OTCace; 1.
Pfam; PF02739; OTCace; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW the EUropean Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by ancentities requires a license agreement (See http://www.isb-sentities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Complete genomic sequence of Corynebacterium glutamicum ATO Submitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
-!- CATALYTIC ACTIVITY: Carbamoyl phosphate + L-aspartate =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TIGREAMS; TIGRO0670; asp_carb_tr; 1.
PROSITE; PS00097; CARBAMOYLTRANSFERASE; 1.
Pyrimidine biosynthesis; Transferase; Comp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
STRAIN=ATCC 13032 /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Corynebacterium glutamicum (Brevibacterium flavum).
Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aspartate carbamoyltransferase (EC transcarbamylase) (ATCase).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QBNQ38;
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                          Pfam; PF00185; OTCace; 1.
Pfam; PF02729; OTCace N; 1.
PRINTS; PR00100; AOTCASE.
                                                                                                                                                                                                                        EMBL; AP005279; BAB99005.1; -.
                                                                                                                                                                                                                                                                            or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nakagawa S.;
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(Rel. 41, Last sequence update)
(Rel. 42, Last annotation update)
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61.1%;
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Pred. No.
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2.1.3.2) (/
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databases.
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RA Strausberg R.D., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Alteschul S.F., Zeeberg B., Buettow K.H., Schaefer C.F., Bhat N.K.,
RA Alteschul S.F., Zeeberg B., Buettow K.H., Schaefer C.F., Bhat N.K.,
RA Alteschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Boak S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Robistein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Robastein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Robastein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Richards S., McKewan P.J., McKernan K.J., Abramson R.D., Mullahy S.J.,
RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
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P49746; Q8WV34;
01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence up
10-0CT-2003 (Rel. 42, Last annotation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE OF 1-736 FROM N.A.
MEDLINE=97477796; PubMed=9331372;
Winfield S.L., Tayebi N., Martin B.N.
"Identification of three additional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Lung;
MEDLINE=96044440; PubMed=7558000;
Adolph K.W., Long G.L., Winfield
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS00097; CARBAMOYLIRANSFERASE; Pyrimidine biosynthesis; Transferase; CG SEQUENCE 312 AA; 33894 MW; F33154F41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Structure and organization of the human thrombospondin 3 gene (THBS3).";
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THBS3 OR TSP3.
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4 MW; F33154F4D033CF22
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       Pfam; PF02412; tsp 3; 13.

Pfam; PF02735; TSPC; 1.

Pfam; PF02735; TSPC; 1.

Pfam; PF02736; TSPC; 1.

SMART; SM00179; EGF CA; 1.

SMART; SM00210; TSPN; 1.

SMART; SM00210; TSPN; 1.

PROSITE; PS01022; EGF 1; FALSE_NEG.

PROSITE; PS01186; EGF 2; 1.

PROSITE; PS01187; EGF CA; 2.

Glycoprotein; Cell adhesion; Calcium-
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MIM; 188062; -.
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EMBL; AF023368; AAC51818.1; -
EMBL; BC018786; AAH18786.1;
PIR; A57121; A57121.
HSSP; P35555; IENN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  domain; Signal.
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107160; P:cell-matrix adhesion; TAS.
1PR001881; EGF_Ca.
IPR006209; EGF_like.
IPR003367; tsp_3.
IPR008859; TSPC.
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                                                                                                                                                                                                                                                                                                                         MEDLINE-22550857; PubMedd-12663927;
Paulsen I.T., Banerjei L., Myers G.S.A., Nelson K.E., Seshadri R.,
Paulsen I.T., Banerjei L., Myers G.S.A., Heidelberg J.F.,
Read T.D., Fouts D.E., Eisen J.A., Gill S.R., Heidelberg J.F.,
Tettelin H., Dodson R.J., Umayam L., Brinkac L., Beanan M.,
Daugherty S., DeBoy R.T., Durkin S., Kolonay J., Madupu R., Nelson W
Vamathevan J., Tran B., Upton J., Hansen T., Shetty J., Khouri H.,
Utterback T., Radune D., Ketchum K.A., Dougherty B.A., Fraser C.M.;
"Role of mobile DNA in the evolution of vancomycin-resistant
Enterococcus faecalis.";
Science 299:2071-2074(2003).

"In SIMILARITY: Belongs to the tautomerase family.
                                                                                               INIT MET
ACT SITE
SEQUENCE
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                                                                                                                                                                   TIGR; EF2859; -.
HAMAP; MF_00718; -; 1.
InterPro; IPR004370; T
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
STRAIN=V583 / ATCC 700802;
                                                                                                                                                                                           EMBL; AE016956; AA082551.1; TIGR; EF2859; -.
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Bacteria; Firmicutes; Lactobacillales; Enterococcaceae; Enterococcus.
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                                                                                                                                                                                                                               or send an email to license@isb-sib.ch).
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                       VEILEGRIECVLSNL 17
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RESULT 11 RS28_KLULA

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P33285;
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between the Swiss Institute or BIOLING THE European Bioinformatics Institute. The European Bioinformatics Institutions as long
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Kluyveromyces.
                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                  gene in Kluyveromyces.";
Yeast 8:949-959(1992).
                                                                                                         "Structure and expression of the
                                                                                                                                       STRAIN=CBS 6556; MEDLINE=93127729; PubMed=1481571;
                                                                                                                                                                                                           Kluyveromyces marxianus (Yeast) (Kluyveromyces fragilis).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Kluyveromyces.
                                                                                                                                                                                                                                                                   40S ribosomal
                                                                                                                                                                                                                                                                                                                                                      KLUMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribosomal protein.
SEQUENCE 67 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam; PF01200; Ribosomal_S28e; 1.
ProDom; PD005541; Ribosomal_S28e; 1.
PROSITE; PS00961; RIBOSOMAL_S28E; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PIR; S30005; S30005.
InterPro; IPR000289; Ribosomal_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; X69582;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE=93127729; PubMed=1481571;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=28985;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           40S ribosomal
                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                      RPS28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SIMILARITY: Belongs to the S28E family of ribosomal proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yeast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hoekstra R., Ferreira P.M.,
                                                                                                                                                                                             NCBI_TaxID=4911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Structure and expression of the ABF1-regulated ribosomal protein

    -:- SIMILARITY: Belongs to the

                                                                                                                                                                                                                                                                                                                                       KLUMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          in Kluyveromyces.";
8:949-959(1992).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
9; Conserv
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                                                                                                                                                                                                                                                               (Rel. 28, Created)
(Rel. 28, Last sequence update)
(Rel. 40, Last annotation update)
al protein S28 (S33).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (Rel. 28, Created)
(Rel. 28, Last sequence update)
(Rel. 37, Last annotation updat)
il protein $28 ($33).
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                                                                                                             Bootsman T.C., Mager W.H., Plant the ABF1-regulated ribosomal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 46;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E5F148312B08A131 CRC64;
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                                                                     family of ribosomal proteins.
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RESULT 13
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Best Local S
Matches 9
                                                                                                                                                                                                 SPECIES-M.tuberculosis; STRAIN-CDC 1551 / Oshkosh;

MEDLINE-22206494; PubMed=12218036;

Fleischmann R.D., Alland D., Eisen J.A., Carpenter L., White O.,

Peterson J., DeBoy R., Dodson R., Gwinn M., Haft D., Hickey E.,

Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M., Salzberg S.L.,

Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M., Salzberg S.L.,

Bolcher A., Utterback T., Weidman J., Khouri H., Gill J., Mikula A.,

Bishai W., Jacobs W.R. Jr., Venter J.C., Fraser C.M.;

"Whole-genome comparison of Mycobacterium tuberculosis clinical and
laboratory strains.";
                                                                                                                                                                                                                                                                                                                                                                                                                       SPECIES=M.tuberculosis; STRAIN=H37Rv;
MEDLINE-98295987; PubMed=9634230;
Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harris Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Gordon S.V., Basham D., Brown D., Chillingworth T., Connor R., Davies R., Devlin K., Feltwell T., Gentles S., Hamlin N., Holroyd Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L., Oliver S., Osborne J., Quail M.A., Rajandream M.A., Rogers J., Rutter S., Seeger K., Skelton S., Squares S., Squares R., Sulston J.E., Taylor K., Whitehead S., Barrell B.G.;
"Deciphering the biology of Mycobacterium tuberculosis from the complete genome sequence.";
Garnier T., Eiglmeier K., Camus J.-C., Medina N., Mansoor H., Pryor M., Duthoy S., Grondin S., Lacroix C., Monsempe C., Simon S. Harris B., Atkin R., Doggett J., Mayes R., Keating L., Wheeler P.1 Parkhill J., Barrell B.G., Cole S.T., Gordon S.V., Hewinson R.G.; "The complete genome sequence of Mycobacterium bovis."; Proc. Natl. Acad. Sci. U.S.A. 100:7877-7882(2003).
                                                                                                                                                                                                                                                                                                                                                                                              Nature
[2]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Putative cytochrome P450 128 (EC 1.14.-.-).
CYP128 OR RV2268C OR MT2330 OR MTCY339.42 OR MB2291C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribosomal protein.
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ProDom; PD005541; Ribosomal_S28e; 1.
PROSITE; PS00961; RIBOSOMAL_S28E; 1.
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PIR; S30006; S30006.
Interpro; IPR000289; Ribosomal S28e.
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Q59572;
                                                                                                            SPECIES=M.bovis; STRAIN=AF2122/97; MEDLINE=22709107; PubMed=12788972;
                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A
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                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mycobacterium tuberculosis, and
                                                                                                                                                                                   Bacteriol. 184:5479-5490(2002).
                                                                                                                                                                                                                                                                                                                                                                                                               393:537-544 (1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  d and this statement is not removed. Usage by and for commercial s requires a license agreement (See http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
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9; Conserv
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Pred. No.
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RESULT 14
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RX MEDILINE-21848401; PubMed=11859360;

RX MEDILINE-21848401; PubMed=11859360;

RA Sgouros J., Peat N., Hayles J., Basham D., Bowman S.,

RA Sgouros J., Peat N., Hayles J., Basham D., Bowman S.,

RA Brooks K., Brown D., Brown S., Chillingworth T., Churcher C.M.,

RA Gentles S., Goble A., Hamlin N., Harris D., Hidalgo J., Hodgson G.,

RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., McLean J.,

RA Holroyd S., Mornsby T., Howarth S., Huckle E.J., Hunt S., McLean J.,

RA Money P., Moule S., Mungall K., Murphy L., Niblett D., Odell C.,

RA Money P., Moule S., Mungall K., Murphy L., Niblett D., Odell C.,

RA Oliver K., O'Neil S., Pearson D., Quail M.A., Rabbinowitsch E.,

RA Autherford K., Rutter S., Saunders D., Seeger K., Sharp S.,

RA Skelton J., Simmonds M., Squares R., Squares S., Stevens K.,

RA Moodward J., Volckaert G., Aert R., Robben J., Grymonprez B.,

RA Modward J., Volckaert G., Reiger M., Schaefer M., Mueller-Auer S.,

RA Meltjens I., Vanstreels E., Rieger M., Schaefer M., Mueller-Auer S.,

RA Gabel C., Fuchs M., Fritzc C., Holzer E., Moestl D., Hilbert H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                             10-OCT-2003 (Rel. 42, Created)
10-OCT-2003 (Rel. 42, Last sequence up
10-OCT-2003 (Rel. 42, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
CENP-A multicopy suppressor protein 2
AMS2 OR SPCC290.04 OR SPCC4F11.01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a c between the Swiss Institute of Bioinformatics and the EWBL the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fo entities requires a license agreement (See http://www.isb-sib.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AMS2_SCHPO
Q9URT4; Q9P7Z1;
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PIR; H70729; H70729.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 458
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MT2330; -.
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POTENTIAL.

IRON (HEME AXIAL LIGAND) (BY SIMILARITY).

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EMBL; AL117389; CAB55767.2; -.
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GO; GO:0005694; C:chromosome; ISS.
GO; GO:0007649; D:cell Cycle; ISS.
GO; GO:0007049; D:cell Cycle; ISS.
InterPro; IPR000679; Znf_GATA.
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P34434;
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SEQUENCE
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                                                                                           Caenorhabditis elegans.
Eukaryota; Metazoa; Nem
Rhabditidae; Peloderina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=22423783; PubMed=12535531;
Chen E.S., Saltoh S., Yanagida M., Takahashi K.;
"A cell cycle-regulated GATA factor promotes centromeric localization of CENP-A in fission yeast.";
Mol. Cell 11:175-187(2003).
STRAIN=Bristol N2;
MEDLINE=94150718; PubMed=7906398;
                                                                                                                                                                01-FEB-1994 (Rel. 28, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Hypothetical protein F44E2.4 in chromosome III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00344; GATA ZN_FINGER_1; FALSE_NEG. PROSITE; PS50114; GATA_ZN_FINGER_2; 1. Transcription regulation; Nuclear protein; Cen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entities requires a license agreement (Some send an email to license@isb-sib.ch).
                                      SEQUENCE FROM N.A.
                                                                                                                                                   F44E2.4.
                                                                                                                                                                                                                                                                                 CAEEL
                                                                      NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zinc-finger
ZN_FING
                                                                                                                                                                                                                                                                                                                                                                                                                          / Match 39.3%;
Local Similarity 40.9%;
nes 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: Required for proper chromosome segregation via regulation of CENP-A localization to the centromere. SUBUNIT: Interacts with CENP-A. SUBCELLULAR LOCATION: Nuclear; centromeric.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY: Contains 1 GATA-type zinc finger.
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697 AA;
                                                                                           Peloderinae;
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78159 MW;
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                                                                                             Caenorhabditis
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5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 46;
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HSSP; P01130; LAJJ.

WORMPED; F44E2.4; CE00182.

Pfam; PF00057; Ldl recept a

SMART; SM00192; LDLa; 1.
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Bonfield J., Burton J., Connell M., Copeey T., Coopeer J., Coulson A
Craxton M., Dear S., Du Z., Durbin R., Favello A., Fraser A.,
Fulton L., Gardner A., Green P., Hawkins T., Hillier L., Jier M.,
Johnston L., Jones M., Kershaw J., Kirsten J., Laisster N.,
Latreille P., Lightning J., Lloyd C., Mortimore B., O'Callaghan M.,
Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen I
Sins M., Smaldon N., Smith A., Smith M., Sonnhammer E., Staden R.,
Sulston J., Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K.,
Waterston R., Watson A., Weinstock L., Wilkinson-Sproat J.,
Wohldman D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commodified and this statement is not removed. Usage by and for commodified and this statement is not removed.
                                                                                                                                                                                                                                                                                                             Hypothetical SEQUENCE 1
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1 QRVEILEGRTECVLSNLRGRTRY
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Copyright (c) 1993 - 2004 Compugen Ltd.
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ALIGNMENTS

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A; Moderate Lype: DNA A; Residues: 1-17;381-387; LLEQQODA',391-395;504-508 <pen> A; Residues: 1-17;381-387; LLEQQODA',391-395;504-508 <pen> A; Note: sequence modified after extraction from NCBI backbone A; Note: the authors translated the codon GAT for residue 31 as B A; Note: an insert compared to other published sequences is considered by authors as like A; Maouche, L.; Tournamille, C.; Hattab, C.; Boffa, G.; Cartron, J.P.; Chretien, S. Blood 78, 2557-2563, 1991 A; Title: Cloning of the gene encoding the human erythropoietin receptor.</pen></pen>	A;Title: The erythropoietin receptor gene: cloning and identification of multiple transc A;Reference number: A53958; MUID:91372359; PMID:1654273 A;Raccession: A53958 A;Raccession: A53958 A;Malcule type: mRNA A;Residues: 1-508 < EHR> R;Penny, L.A.; Forget, B.G. Genomics 11, 974-980, 1991 Genomics 11, 974-980, 1991 A;Riels: Genomic organization of the human erythropoietin receptor gene. A;Reference number: A55280; MUID:92147143; PMID:1664413 A;Accession: A55280	A;Accession: A49824 A;McLecule type: DNA A;Mclecule type: DNA A;Residues: 1-508 <nog> A;Cross-references: GB:S45332; NID:g255496; PIDN:AAB23271.1; PID:g255497 A;Experimental source: placenta A;Note: sequence extracted from NCBI backbone (NCBIN:113293, NCBIP:113294) R;Bhrenman, K.; St. John, T. R;Bhrenman, K.; St. John, T. Exp. Hematol. 19, 973-977, 1991</nog>	A;Cross-references: GB:M60459; NID:g182244; PIDN:AAA52403.1; PID:g182245 R;Winkelmann, J.C.; Penny, L.A.; Deaven, L.L.; Forget, B.G.; Jenkins, R.B. Blood 76, 24-30, 1990 A;Title: The gene for the human erythropoietin receptor: analysis of the coding sequence A;Reference number: A60160; MUID:90304334; PMID:2163695 A;Accession: A60160 A;Status: not compared with conceptual translation A;Molecule type: mRNA; DNA A;Residues: 1-101, 'R', 103-188, 'RP', 191-243, 'E', 245-508 <win> R;Noguchi, C.T.; Bae, K.S.; Chin, K.; Wada, Y.; Schechter, A.N.; Hankins, W.D. Blood 78, 2548-2556, 1991 A;Title: Cloning of the human erythropoietin receptor gene. A;Reference number: A49824; MUID:92399733; PMID:1668606</win>	RESULT 1 ZUHUR erythropoietin receptor precursor - human C;Species: Homo sapiens (man) C;Date: 12-Feb-1993 #sequence revision 05-Apr-1995 #text change 22-Jun-1999 C;Accession: A43799; A60160; Ā49824; A53958; A55280; I52563 R;Jones, S.S.; D'Andrea, A.D.; Haines, L.L.; Wong, G.G. Blood 76, 31-35, 1990 A;Title: Human erythropoietin receptor: cloning, expression, and biologic characterizati A;Reference number: A43799; MUID:90304340; PMID:2163696 A;Molecule type: mRNA A;Molecule type: mRNA A;Molecule type: mRNA A;Mesidues: 1-508 <jon></jon>

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A;Map position: 19p13.3-19p13.2

A;Introns: 39/1; 84/2; 143/1; 195/3; 247/1; 276/2; 305/3

C;Superfamily: erythropoietin receptor; cytokine receptor homology

C;Keywords: alternative splicing; cytokine receptor; glycoprotein; transmembrane protein

F;1-24/Domain: signal sequence #status predicted <SIG-
F;25-508/Product: erythropoietin receptor #status predicted <MAT>
F;25-250/Domain: extracellular #status predicted <EXT>
F;52-250/Domain: cytokine receptor homology <CRS>
F;23-237/Region: WSXWS motif
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A; Residues: 1-265 < KUR>
R; Laccombe, C.; Chretien, S.; Lemarchandel, V.; Mayeux, P.; Romeo,
J. Biol. Chem. 266, 6952-6956, 1991
J. Title: Spleen focus-forming virus long terminal repeat insertion
A; Title: Spleen focus-forming 191201346; PMID:1849897
A; Reference number: 149653; MUID:91201346; PMID:1849897
A; Recession: 149653
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R;Kuramochi, S.; Ikawa, Y.; Todokoro, K.
J. Mol. Biol. 216, 567-575, 1990
A;Title: Characterization of murine erythropoietin receptor genes.
A;Reference number: S13249; MUID:91080149; PMID:2175360
A;Accession: S14081
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erythropoietin receptor precursor, membrane-bound form - C;Species: Mus musculus (house mouse)
C;Date: 28-Sep-1990 #sequence_revision 05-Apr-1995 #text_
                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: GB:M62360; NID:g193199; PIDN:AAA37582.1; PID:g193200 C;Superfamily: erythropoietin receptor; cytokine receptor homology C;Keywords: cytokine receptor; transmembrane protein F;52-238/Domain: cytokine receptor homology <CRS>
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A; Residues: 1-24 < RES>
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C;Genetics:
                                                                                        A32385
                                                                                                                   RESULT 3
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|Species: Mus musculus (house mouse)
|Species: Mus musculus (house mouse)
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Best Local
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                                                                                                                                                                                                                                                                                                                    Conservative
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#sequence_revision 05-Apr-1995 #text_change 22-Jun-1999
                                                                                                                                                                                                                                                                                                                                              93.2%;
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Pred. No. 2.3e-09;
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A;Residues: 1-507 < CDAA>
A;Cross-references: GB.J04843; NID:g193090; PIDN:AAA37571.1; PID:g309219
A;Experimental source: murine erythroleukemia (MEL) cells, subclone 745
R;Kuramochi, S; Ikawa, Y; Todokoro, K.
J. Mol. Biol. 216, 567-575, 1990
A;Title: Characterization of murine erythropoietin receptor genes.
A;Reference number: S13249; MUID:91080149; PMID:2175360
A;Accession: S13249
                           C;Superfamily: erythropoietin receptor; cytokine receptor homology C;Keywords: cytokine receptor; glycoprotein; transmembrane protein F;1-24/Domain: signal sequence #status predicted <SIG> F;25-507/Product: erythropoietin receptor #status predicted <MAT> F;25-249/Domain: extracellular #status predicted <EXT>
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c;Superfamily: erythropoietin receptor; cytokine receptor homology
C;Keywords: alternative splicing; cytokine receptor; glycoprotein;
F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-207/Product: erythropoietin receptor #status predicted <MAT>
F;25-249/Domain: extracellular #status predicted <EXT>
F;25-238/Domain: cytokine receptor #status cRKS>
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A;Residues: 1-507 <KUR>
A;Crose-references: EMBL:X53081; NID:g50861; PIDN:CAA37248.1; PID:g50862
A;Experimental source: murine erythroleukemia K-1 cells
A;Experimental source:
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A;Title: Expression cloning of the murine erythropoietin receptor.
A;Reference number: A32385; MUID:89195238; PMID:2539263
A;Accession: A32385
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                                                                                                                                                                                                                A;Cross-references: GB:D13566; NID:g286209; PIDN:BAA02761.1; PID:g286210
A;Experimental source: PC12 and erythroid cells
A;Note: sequence extracted from NCBI backbone (NCBIN:132811, NCBIP:132813)
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A;Molecule type: mRNA
A;Residues: 1-507 <MAS>
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R;Masuda, S.; Nagao, M.; Takahata, K.; Konishi, Y.; Gallyas Jr J. Biol. Chem. 268, 11208-11216, 1993
J. Biol. Chem. 268, 11208-11216, 1993
A;Title: Functional erythropoietin receptor of the cells with A;Reference number: A46713; MUID:93266574; PMID:7684373
A;Accession: A46713
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R;Hino, M.; Tojo, A.; Misawa, Y.; Morii, H.; Takaku, F.; Shibuya, M.
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;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 16-Jun-2000
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cytokine
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91.3%;
receptor homology
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Pred. No. 4.6e-09;
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A,Cross-references: GB:L38969; NID:g886298; PIDN:AAC41762.1; PID:g886299; GB:L38970 R;Winfield, S.L.; Tayebi, N.; Martin, B.M.; Ginns, E.I.; Sidransky, E. Genome Res. 7, 1020-1026, 1997 A;Title: Identification of three additional genes contiguous to the glucocerebrosidase A;Reference number: Z16482; MUID:97474796; PMID:9331372 A;Reterence number: Z16482; MUID:97474796; PMID:9331372 A;Accession: T08830 A;Status: translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-736 <WIN>
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C;Date: 03-Nov-1995 #sequence_revision 03-Nov-1995 #text_change 21-Jan-2000
C;Accession: A57121; T08830
R;Adolph, K.W.; Long, G.L.; Winfield, S.; Ginns, E.I.; Bornstein, P.
Genomics 27, 329-336, 1995
A;Title: Structure and organization of the human thrombospondin 3 gene (THBS3).
A;Reference number: A57121; MUID:96044440; PMID:7558000
A;Accession: A57121
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-956 ADO>
A;Residues: 1-956 ADO>
A;Residues: 1-956 ADO>
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                                                                                                                                                                                                                                                                                                                                                         RESULT
S30006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A,Description: mediates cell-to-matrix and cell-to-cell interactions C;Superfamily: thrombospondin 3, EGF homology F,374-412/Domain: EGF homology <EGF>
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A;Introns: 27/1; 96/1; 181/3; 216/1; 225/1; 256/1; 270/1; 319/3; 366/3; 392/3; 443/3; 48
C;Function:
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                                                                                                                                Yeast 8, 949-959, 1992
A;Title: Structure and expression of the ABF1-regulated ribosomal protein S33
A;Reference number: S30005; MUID:93127729; PMID:1481571
                                                                                                                                                                                                                                                                              ribosomal protein S28.e - yeast (Kluyveromyces marxianus) N;Alternate names: ribosomal protein YS33 C;Species: Kluyveromyces marxianus
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A;Cross-references: GDB:409953; OMIM:188062
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                                          A; Cross-references: GB:S52656
                                                                 A; Molecule type: DNA
A; Residues: 1-67 < HOE>
                                                                                                          A; Accession: S30006
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Pred. No. 4.6e-09;
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RESULT 8
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C;Species: Kluyveromyces marxianus var. lactis, Candida sphaerica
C;Date: 07-May-1993 #sequence_revision 07-May-1993 #text_change 07-May-1999
C;Accession: S3005
R;Hoekstra, R.; Perreira, P.M.; Bootsman, T.C.; Mager, W.H.; Planta, R.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Gene: S33
C;Superfamily: rat ribosomal protein S28
C;Keywords: protein biosynthesis; riboso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Title: Structure and expression of the ABF1-regulated ribosomal protein S33 gene in A;Reference number: S30005; MUID:93127729; PMID:1481571 A;Accession: S30005
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                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-288 < SI
                                                                                                                                                                                                                                                                                                                                                                  R;Senkevich, T.G.; Bugert,
Science 273, 813-816, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                             probable DNA-binding protein 46L - Molluscum contagiosum virus N,Alternate names: MC046L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: GB:S53420 C;Genetics:
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A; Residues: 1-67 < HOE>
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                                                                                                                                                                                                                                                                                                          A; Accession: T30648
                                                                                                                                                                                                                                                                                                                          A;Title: Genome sequence of a human tumorigenic poxvirus: Prediction of specific host A;Reference number: Z20876; MUID:96325459; PMID:8670425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yeast 8, 949-959, 1992
                                                                                                                                                                                         A; Note: MC046L
                                                                                                                                                                                                                             A;Cross-references: EMBL:U60315; NID:g1491943; PIDN:AAC55174.1; PID:g1491989
                                                                                                                                                                                                                                                                                     A; Status: preliminary; translated
                                                                                                                                                                                                                                                                                                                                                                                                      Species: Molluscum contagiosum virus 1
,Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 27-Oct-2003
,Accession: T30648
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8; Conserv
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ECIYTNLRGGTR 144
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Pred. No. 3.
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Pred. No. 17;
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N;Contains: oxidoreductase (b. 1....)

R;Contains: oxidoreductase (b. 1....)

C;Species: Mycobacterium tuberculosis

C;Dacte: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 21-Jul-2000

C;Accession: H70729

R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.

R;Cole, S.T.; Brosch, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.;

; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Squares, Squar
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A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-710 <BAR>
A;Cross-references: EMBL:AL117389; PIDN:CAB55767.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable transcription regulator - fission yeast (Schizosaccharomyces pombe) C;Species: Schizosaccharomyces pombe C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 21-Jan-2000 C;Accession: T41352; T38826 R;Barrell, B.G.; Rajandream, M.A.; Quail, M.; Harris, D.; Lyne, M.
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A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete A;Reference number: A70500; MUID:98295987; PMID:9634230
A;Accession: H70729
A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA
A;Residues: 1-489 <COL>
A;Cross-references: GB:Z77163; GB:AL123456; NID:g3261610; PIDN:CAB00967.1; PID:g1
A;Experimental source: strain H37Rv
C;Genetics:
                                                                                                                                                                                          F44E2.4 protein - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 20.Feb-1995 #sequence_revision 2 C;Accession: S44821
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C;Keywords: oxidoreductase
F;322-457/Domain: cytochrome P450 homology <P45>
                               A; Reference number: S44816
A; Accession: S44821
                                                                                                submitted to the EMBL Data A; Description: Sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Barrell, B.G.; Rajandream, M.A.; submitted to the EMBL Data Library,
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A;Status:
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ibrary, June 1999
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A;Molecule type: DNA
A;Residues: 1-1609 <AND>
A;Residues: 1-1609 <AND>
A;Crose-references: EMBL:L23646; NID:g388595; PID:g388601
C;Genetics: 107/2; 173/2; 228/3; 1165/2; 1216/3; 1231/3; 1258/2; 1300/2; 1321/3; 1:C;Superfamily: LDL receptor ligand-binding repeat homology
F;11-45/Domain: LDL receptor ligand-binding repeat homology <LDL>
                                                                                                                                                                                                                                                                                C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 28-Jul-2003 C;Accession: D83406  
R;Stover, C:K:; Pham, X:Q:; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hich adman, S:; Yuan, Y:, Brody, L.L.; Coulter, S.N.; Folger, K:R.; Kas, A.; Lark:; Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 13
D83406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: GB:D16247; NID:g466572; PIDN:E C;Superfamily: tobacco ATP-dependent RNA helicase C;Keywords: ATP; nucleotide binding; P-loop F;18-56/Domain: MW repeat homology <WWl>F;189-196/Region: nucleotide-binding motif A (P-lo F;294-299/Region: nucleotide-binding motif B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATP-dependent RNA helicase DB10 - wood tobacco C;Species: Nicotiana sylvestris (wood tobacco) C;Date: 10-Sep-1999 #sequence_revision 10-Sep-194639
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S42639
                                                                                                                                                                                                                           A;Title: Complete genome sequence of Pseudomonas aeruginosa A;Reference number: A82950; MUID:20437337; PMID:10984043 A;Accession: D83406
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A; Residues: 1-607 < ITA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Structure and expression of a cDNA encoding an A;Reference number: $42639; MUID:94154240; PMID:8111024 A;Accession: $42639 A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R; Itadani, H.; Su
Plant Mol. Biol.
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                                                                                                                                                                                                                                                                                                                                                                                              probable transmembrane sensor PA1911 [imported] - C;Species: Pseudomonas aeruginosa
                                                                                                               C; Genetics:
                                                                                                                                 A; Experimental
                                                                                                                                                     A;Cross-references:
                                                                                                                                                                      A; Residues: 1-316 <STO>
                                                                                                                                                                                        A; Molecule type: DNA
                                                                                                                                                                                                        A; Status: preliminary
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Best Local S
Matches 8
                 Query Match
Best Local
                                                                          Gene: PA1911
Superfamily:
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12; Conser
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8; Conserv
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                                                                          Fe2+-dicitrate sensor, transmembrane
                                                                                                                                   source: strain
38.5%;
llarity 54.5%;
Conservative
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                                                                                                                                   PAO1
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Pred. No. 44;
4; Mismatches
                 Score 45;
Pred. No.
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Pred. No.
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licase DB10; WW repeat homolog
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1e+02;
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                                     DB 2;
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                                       Length 316
                                                                            component
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A.; Larbig,
   Gaps
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K.; Lim,
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N;Alternate names: neural cell adhesion molecule AxCAM
C;Speciaes: Gallus gallus (chicken)
C;Date: 30-Sep-1993 #sequence revision 30-Sep-1993 #text_change 21-Jan-2000
C;Accession: S22383; S34107; S69332; S22128
R;Zuellig, R.A.; Rader, C.; Schroeder, A.; Kalousek, M.B.; von Bohlen und Halbach, F Bur. J. Biochem. 204, 453-463, 1992
A;Title: The axonally secreted cell adhesion molecule, axonin-1. Primary structure, A;Reference number: S22383; MUID:92174898; PMID:1311675
A;Accession: S22383
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(;Species: Mycobacterium tuberculosis
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 21-Jul-2000
C;Accession: D70506
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, S.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroyd, S. Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, S37-S44, 1998
A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genome A;Reference number: A70500; MUID:98295987; PMID:9634230
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D70506
A;Cross-references: EMBL:X79607
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1994
C;Superfamily: contactin; fibronectin type III repeat homology; immunoglobulin homology
C;Keywords: cell adhesion
F;1-23/Domain: signal sequence #status predicted <SIG>
F;1-23/Domain: immunoglobulin homology <IMM>
F;336-392/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: protein
A;Residues: 29-49,51-80,84-95,100-117;120-128,130-141;143-176;243-254;256-296;303-336;33R;Giger, R.J.; Vogt, L.; Zuellig, R.A.; Rader, C.; Henehan-Beatty, A.; Wolfer, D.P.; Sor Eur. J. Biochem. 227, 617-628, 1995
A;Title: The gene of chicken axonin-1. Complete structure and analysis of the promoter.
A;Reference number: S69332; MUID:95172044; PMID:7867620
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A; Residues: 1002-1036 <GIG>
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A; Residues: 1-1036 < ZUE1>
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C;Superfamily: conserved hypothetical protein b0835
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A; Residues: 1-512 < COL>
                                                                                                                                                                                                                                                                                                                                             A; Accession: S69332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 38.5%;
Local Similarity 42.3%;
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Query Match

38.5%;

Score 45;

DB 2;

Length 1036;

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Best Local Similarity 47.1%; Pred. No. 92;
Matches 8; Conservative 2; Mismatches 7; Indels 0; G

Qy 7 EGRTECVLSNLRGRTRY 23
Db 296 EGTYECEAENIKGRDTY 312
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Search completed: May 6, 2004, 12:47:27 Job time : 22 secs



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OM protein - protein search, using sw model
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Perfect score:
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Listing first 45 summaries
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A Geneseq_29Jan04:*
1: geneseqp1980s:*
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117
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                            1586107 seqs, 282547505 residues
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	. 10	9	8	7	D	տ	4	ω	N	ъ	Result No.
52	102	109	109	109	109	111	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	117	Score
44.4	87.2	93.2	93.2	93.2	93.2	94.9	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	Query Match I
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Claim 8; Page This sequence activation se Activation se Activation se are separate important in internalisati Use of pepti can retard on stabilising t surface. This In addition, meaning that ligand, causi probably brou	WPI; 1999-5 Modulating of, e.g. ok	Hómo sapic WO99421277 26-AUG-19: 23-FEB-19: 24-FEB-19: (RECE-) RI	LT 1 9411 AAY394 AAY394 AAY394 Human Human	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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sh. ptide derived from the man erythropoletin (Ep) lived in modulation of binding site. Activations: in the modulation of activation of the activation of activation of activation of active recepor internalisation, the number of active recepof increasing signalliclassed as a type 2 cen actually replace the naturally replace the naturally replace the indent activation. This	-2 cell surface receptors	10. 38.	ALIGNMENTS ptide; 23 AA. ntry) receptor-derived activation peptide. receptor; activation; internalisation	AAW29667 AAB84225 AAE13249 ABG13674 ABP63791 AAY02136 AAB97378 AAB66034 AAB66063 AAB66063 AAB66063 AAB032717 AAB66063 AB032717 AAB66033
extracellular No) receptor responses and on sequences are freeptor receptor receptor receptor receptor receptor in a receptor reby increasing or the cell form on the cell flaund. Ill surface receptor, requirement for the activation is in which one peptide	used in treatment		tide.	Aaw29667 Homo sapi Aab84225 A tumour Aae13249 Human tum Abg13674 Novel hum Abg65639 Bifidobac Abp43791 Human kri Aay02136 Protein e Aab66034 Human TAN Abo32694 Secreted Aab84220 A human tum Aab66063 Murine pr Aab632717 Secreted Aab86033 Human TAN Abo32693 Secreted Aab86033 Human TAN Abo32693 Secreted Aab66083 Murine TAN Abo32693 Secreted Aay02135 HTHB247 p Abg76512 DNA encod Aab66083 Murine TA

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This invention corresponds to a novel composition comprising any of 34 cp peptides consisting of activation sequences from the extracellular commology searching with a peptide sequence from the MRC class I alpha I domain and correspond to the activation sequences of the associated receptor. Activation sequences are involved in the internalisation of receptors and so these peptides may be used to modulate the internalisation and or activation of these receptors. The peptides of the invention may also be used to increase effect of ligand signalling in type 2 diabetes and can replace the ligand normally required for activation. This is useful when the ligand is a hormone and is difficult to obtain. The peptides may also interact syneryistically with the ligand or function as antagonists of receptor signalling. The peptides can also be used in drug screening to identify compounds that modulate receptor internalisation or function as ligand replacements. Use of these peptides can increase the effect of therapeutic hormones (optionally combined that receptors from which they derive. The present sequence represents the human contains the human and the present sequence represents the human contains and contains and contains the human contains and contains 
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Best Local S
Matches 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Composition comprising peptides from receptor extracellular domains, useful for e.g. modulating receptor internalization and activation such as increasing insulin activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Olsson L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-AUG-1996;
23-JAN-1997;
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                            Erythropoietin receptor
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96US-00701382.
97US-00788820.
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                          activation sequence oligopeptide of the invention
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Pred. No. 3.6e-11;
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                                            The present sequence is that of an erythropoietin (EPO) mimetic peptide which has no homology to natural EPO but has similar biological properties in that it activates the EPO receptor acting as an agonist. The EPO mimetic peptide can be included in novel fusion proteins of the invention also including a transferrin (Tf. see ABP72820) molety engineered to extend the serum half-life or bioavailability. The EPO mimetic peptide can be fused to the N- or C-terminus of Tf. or inserted into, or used to replace part of Tf. such that the Tf acquires EPO activity. Modified Tf fusion proteins of the invention can be used in the diagnosis, prognesis, prevention and/or treatment of diseases and/or disorders of the endocrine, nervous, immune, respiratory, cardiovascular, reproductive and digestive systems, diseases and/or disorders relating to the blood or to cell proliferation, inflammatory conditions and infectious diseases, or to deliver a therapeutic agent to a cell or
                                                                                                                                                                                                                                                                                                                                                                       New fusion protein, useful in the diagnosis and treatment of diseases of disorders relating to the respiratory, cardiovascular and digestive systems, comprises a transferrin protein fused to a therapeutic protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fungicide; hepatotropic; antirheumatic; antiarthritic; antigout; tranquillizer; vulnerary; antidiabetic; nephrotropic; antipyretic; gastrointestinal; gene therapy; transgenic animal; erythropoietin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transferrin; neuroprotective; cerebroprotective; vasotropic; antiparkinsonian; nootropic; anti-HIV; antiasthmatic; antiallergic; cytostatic; immunosuppressive; antiatherosclerotic; cardiant;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-332916/31
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30-NOV-2001; 2001US-0334059P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gynaecological; immunostimulant; antianaemic; haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Erythropoietin mimetic peptide.
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                                   blood-brain barrier
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Sequence

23

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Length 23;

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RESULT 4
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                                                                                                                                                                           Matches
                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                   contg. the antibody may be used in the diagnosis and treatment of patients having disorders associated with low red blood cell levels, e. anaemia. The antibodies are also useful in methods and kits for detecti EPORs in biological in biological samples
                                                                                                                                                                                                                                                                                                     erythropoiesis by stimulating the proliferation and/or differentiation of erythroid progenitor cells to erythrocytes. Pharmaceutical compsns.
                                                                                                                                                                                                                                                                                                                             AAR89960-R89965 and AAR98936-R98939 are overlapping, synthetic human erythropoietin receptor (sHuEPOR) peptides which span residues 1 to 244 of the human EPOR. The peptides are used to map the EPOR binding epitope of an EPOR monoclonal antibody which binds to EPORs and stimulates
                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis and treatment of pred blood cell levels, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR98938;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR98938 standard;
                                                                                                                                                                                                                            Sequence 30 AA;
                                                                                                                                                                                                                                                                                                                                                                                              Example 6; Page 34; 61pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Monoclonal antibodies stimulating an erythropoietin receptor - useful in diagnosis and treatment of patients having disorders associated with low
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1996-117004/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic human erythropoietin receptor peptide,
              21-DEC-2000
                                      AAB21686;
                                                              AAB21686 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monoclonal antibody; erythropoietin receptor; diagnosis; anaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-SEP-1996
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                                                                                                                                                                           23;
                                                                                                                                                  1 ORVEILEGRIECVLSNLRGRIRY 23
                                                                                                                                                                                       Similarity
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                                                                                                                           QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QRVEILEGRIECVLSNIRGRIRY 23
                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94US-00280864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95WO-US009458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            erythrocyte production; epitope mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide;
                                                              peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.08; 27
                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30
                                                                                                                                                                                                                                                                                                                                                                                                                       anaemia.
                                                              211 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 3.6; Mismatches
                                                                                                                                                                                      Score 117; DB 2;
Pred. No. 4.9e-11;
                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.6e-11;
hes 0;
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                                                                                                                                                                                                  Length 30;
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                                                                                                                                                                                                                                                                                          RESULT 6
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method for screening for a ligand analog, comprising adding a candidate ligand to a non-naturally occurring cell surface receptor analog e.g. erythropoietin receptor (EPOR), and determining the binding of the ligand to the analog. The present sequence is a mature human erythropoietin receptor (EPOR) extracellular domain. Protein Design Automation was carried out on the present sequence, so that it may be used in the present invention as a cell surface receptor analog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Screening for ligand analogs and agents which modulate ligand-receptor binding, comprises adding a test ligand to a non-naturally occurring constraint exceptor analog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mature erythropoietin receptor EPOR extracellular domain
                                    11-FEB-1999;
29-APR-1999;
                                                                                                                      WO200047612-A2
                                                                                                                                             Homo sapiens.
                                                                                                                                                                                          Human mature erythropoietin receptor EPOR extracellular domain
                                                                                                                                                                                                                    21-DEC-2000
                                                                                                                                                                                                                                            AAB21685;
                                                                                                                                                                                                                                                                  AAB21685 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 211 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      analog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Fig 8; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-549135/50
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29-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ligand; cell surface receptor; erythropoietin; EPOR; human;
                                                                       11-FEB-2000; 2000WO-US003665.
                                                                                                                                                                  Ligand; cell
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                                                                                               17-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                               QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                ORVEILEGRIECVLSNLRGRIRY 183
                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                  surface receptor; erythropoietin; EPOR; human
                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0120009P
99US-0131674P
                                    99US-0120009P
                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 117; DB 3; 100.0%; Pred. No. 4.7e-10; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                    225
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                                                                                                                                                                                                                                                                                                                                                                                                      Length 211;
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(XENC-) XENCOR INC

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RESULT 7
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Best Local :
                 The present sequence is truncated human EpoR (erythropoietin receptor). Transfection of constitutively active EpoR (1439) by electroporation into a cytokine-dependent cell line supports cell population expansion in the absence of exogenous cytokines. Mutant human EpoR is used in treatment of disorders related to inadequate EpoR signalling. The transfected cells
                                                                                                                                                                                                                     used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Truncated human EpoR; erythropoietin receptor; hypersensitive EpoR(t439); mutant human EpoR; EpoR signalling; cancer; infectious disease; HIV; sickle cell anaemia; cytostatic; antimicrobial; antiviral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell surface receptor analog e.g. erythropoietin receptor (BPOR), and determining the binding of the ligand to the analog. The present sequence is a mature human erythropoietin receptor (BPOR) extracellular domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Screening for ligand analogs and binding, comprises adding a test surface receptor analog.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a method for screening for a ligand analog, comprising adding a candidate ligand to a non-naturally occurring
                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                               P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                      Bell D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY44622;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 225 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunostimulant;
                                                                                                                                                                                                                                                                                                                                                                                                                       (HEMO-)
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                                                                                                                                                                                                                                                                                                                      2000-136979/12.
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                                                                                                                                                                                                                                               free defined medium useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23;
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                                                                                                                                                                                                                                                                                                                                                                                                                       HEMOSOL INC
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                                                                                                                                                                                                                                                                                                                                                                      Matthews KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QRVEILEGRIECVLSNLRGRIRY 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   analog
                                                                                                                                                                                                                     production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human
                                                                                                                                                                  Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fig 8; 82pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98CA-02241576.
99CA-02260332.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                anti-anaemic
                                                                                                                                                                  61pp;
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                                                                                                                                                                                                                        of.
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therapy to
                                                                                                                                                                                                                     hemoglobin
                                                                                                                                                                                                                                                                                                                                                                      Mueller
                                                                                                                                                                        English.
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Pred. No. 5e-10;
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                                                                                                                                                                                                                                                 efficient
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QRVEILEGRTECVLSNLRGRTRY 23

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RESULT 8
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Best Local
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                                                                                                                                                                                                   The present sequence is R154C truncated human EpoR (erythropoietin receptor). Transfection of constitutively active EpoR (£439; R154C) electroporation into a cytokine-dependent cell line supports cell population expansion in the absence of exogenous cytokines. Mutant EpoR is used in treatment of disorders related to inadequate EpoR
                                                                                                                                   EpoR is used in treatment of disorders related to inadequate EpoR signalling. The transfected cells may also used in gene therapy to cancer, infectious diseases (e.g. HIV), sickle cell anaemia, and conditions related to abnormal expression of erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                              used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Truncated human EpoR; erythropoietin receptor; hypersensitive EpoR(t439); mutant human EpoR; EpoR signalling; cancer; infectious disease; HIV; sickle cell anaemia; cytostatic; antimicrobial; antiviral;
                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                      Serum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bell D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-JUN-1998;
25-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R154C truncated human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-2000
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                                                                                                                                                                                                                                                                                                                                                Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9967360-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunostimulant;
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erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                              for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000-136979/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   194
                                                                                                                                                                                                                                                                                                                                                                                           free defined medium useful for production of hemoglobin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23;
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                                                                                             438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matthews KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                              Fig 10; 61pp; English.
100.0%; ilarity 100.0%; Conservative 0;
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99CA-02260332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mia; cytostatic; antimicrobial;
anti-anaemic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Wild type Arg substituted by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anaemia,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mueller
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                       Score 117; DB 3;
Pred. No. 1.1e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 117; DB 3;
Pred. No. 1.1e-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ₿
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                                          Length 438;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
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ID ADE28671

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07-DEC-2001; 2001US-0338285P.
12-DEC-2001; 2001US-0341346P.
17-DEC-2001; 2001US-0341477P.
17-DEC-2001; 2001US-0341477P.
27-DEC-2001; 2001US-0342592P.
27-DEC-2001; 2001US-0344297P.
31-DEC-2001; 2001US-0344297P.
31-DEC-2001; 2001US-0344297P.
31-DEC-2001; 2001US-0344293P.
17-MAY-2002; 2002US-033288P.
17-MAY-2002; 2002US-0380981P.
17-MAY-2002; 2002US-0381495P.
28-MAY-2002; 2002US-0383344P.
29-MAY-2002; 2002US-038334P.
29-MAY-2002; 2002US-0383829P.
29-MAY-2002; 2002US-0383829P.
29-MAY-2002; 2002US-0383829P.
29-MAY-2002; 2002US-038653P.
26-AUG-2002; 2002US-0401788P.
31-OCT-2002; 2002US-040533P.
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                                                                                                                                                                                                                                                       Alsobrook JP, Alvarez E, Anderson DW, Baron M, Burgess CE, Casman SJ, Chapoval A, Dhanabal M, Ellerman K, Ettenberg S, Gangolli EA, Gerlach V Grosse WM, Guo X, Hackett C, Ji W, Kekuda R, Lepley DM, Li L, Macdougall JR, Malyankar UM, Mezes PS, Miler CE, Millet I, Mishra VS, Padi Pena CEA, Peyman JA, Rastelli L, Rieger DK, SF Smithson G, Starling G, Spytek KA, Stone DJ, T Vernet CAM, Zerhusen BD, Zhong M;
                                                      preventing, diagnosing or treating NOVX-as osteoarthritis, obesity, atherosclerosis,
                                                                                  New isolated NOVX polypeptides and preventing, diagnosing or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-NOV-2001;
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                              asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9
                                                                                                                                                                                                            2003-441555/41.
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                                 or intections
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                                                         polynucleotides, useful for
NOVX-associated disorders, e.g.
erosis, cancer, Parkinson's disease,
                                                                                                                                                                                                                                                                                     rlach VL, Gorman L;
da R, Khramtsov NV;
r UM, Mazur A, Mcqu
, Padigaru M, Pattu
DK, Shenoy SG, Shim
DJ, Tchernev VT, T
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Edinger
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                                                                                                                                                                                                                                                                                                                       Patturajan
, Shimkets I
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                                                                                                                                                                                                                                                                                                                                                                                 Mcqueeney
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RESULT 10
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Best Local S
Matches 23
05-NOV-2001;

05-DEC-2001;

07-DEC-2001;

12-DEC-2001;

17-DEC-2001;

17-DEC-2001;

20-DEC-2001;

27-DEC-2001;

31-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptides, nucleic acid molecules and antibodies may be useful fureating or diagnosing diseases including metabolic disorders such diabetes and obesity, infectious diseases, anorexia, cancer, cardiovascular diseases including hypertension and atherosclerosis, neurodegenerative disorders such as Alzheimer's disease, Parkinson' disease, and anilony impure discreta such as Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nootropic; neuroprotective; antiparkinsonian; anticonvulsant; osteopathic; antiarthritic; antiinflammatory; dermatological; antiasthmatic; antiilipaemic; metabolic; diabetes; obesity; infectious; antorexia; cancer; cardiovascular; hypertension; atherosclerosis; neurodegenerative; Alzheimer's disease; Parkinson's; epilepsy; immune; osteoparthritis; haemopoietic; inflammatory skin; asthma; dyslipidaemia; neurogenesis; cell differentiation; proliferation; haemopoiesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NOVX; antidiabetic; anorectic; cardiant; hypotensive; antidiateriosclerotic; virucide; antibacterial; fungicide; protozoacide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 458 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cardiant,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human NOV15b protein - SEQ ID 50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JAN-2004
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                                                                                                                                                                                                                                                                                                          05-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                      15-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003040330-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               wound healing; angiogenesis; gene therapy; chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human NOV protein of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QRVEILEGRTECVLSNLRGRTRY 216
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2001US-0338626P.
2001US-0336600P.
2001US-0341346P.
2001US-0341477P.
2001US-0341477P.
2001US-0341540P.
2001US-0342592P.
2001US-0344297P.
2001US-0344903P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                          2002WO-US035536
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Pred. No. 1.1e-09;
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RESULT 11
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AC ADE28
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DT 29-JA
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Best Local (
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15-MAY-2002;
17-MAY-2002;
28-MAY-2002;
28-MAY-2002;
29-MAY-2002;
29-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pena CEA, I
Smithson G,
                                                                                                                                                                                                                                                                                                                           disease and epilepsy, immune disorders e.g. osteoarthritis, haemopoietic disorders, inflammatory skin disorders, asthma and dyslipidaemia. Furthermore, the nucleic acids and polypeptides may also be used to identify molecules that modulate or inhibit neurogenesis, cell differentiation and proliferation, haemopoiesis, wound healing and angiogenesis, as well as in gene therapy. Finally, the nucleic acids may be used as hybridisation probes, in chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The current sequence is that the human NOV protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polypeptides, nucleic acid molecules and antibodies may be useful f
treating or diagnosing diseases including metabolic disorders such
diabetes and obesity, infectious diseases, anorexia, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel isolated NOVX polypeptide. The polypeptide of the invention demonstrates, antidiabetic, anorectic, cardiant, hypotensive, antiarteriosolerotic, virucide, antibacterial, fungicide, protozoacide, nootropic, neuroprotective, antiparkinsonian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preventing, diagosteoarthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cardiovascular diseases including hypertension and atherosclerosis, neurodegenerative disorders such as Alzheimer's disease, Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            anticonvulsant, osteopathic, antiarthritic, antiinflammatory, dermatological, antiasthmatic and antilipaemic activities. The
                  29-JAN-2004
                                                                                 ADE28675 standard;
                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Grosse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mezes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lepley DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       isolated NOVX polypeptides and polynucleotides, useful for venting, diagnosing or treating NOVX-associated disorders, e.g. earthritis, obesity, atherosclerosis, cancer, Parkinson's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2003-441555/41.
DB; ADE28672.
                                                                                                                                                                 194
                                                                                                                                                                                                                               23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURAGEN
                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                458
                                                                                                                                                                                               ORVEILEGRIECVLSNLRGRIRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID
                                                                                                                                                               ORVEILEGRIECVLSNIRGRIRY 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JP, Alvarez E, Anderson DW, Baron M, Boldog FL;
, Casman SJ, Chapoval A, Dhanabal M, Edinger SR,
Casman SJ, Chapoval A, Dhanabal M, Edinger SR,
Casman L;
Guo X, Hackett C, Ji W, Kekuda R, Khramtsov NV;
Li L, Macdougall JR, Malyankar UM, Mazur A, Mcqn
Miller CB, Millet I, Mishra VS, Padigaru M, Patty
Peyman JA, Rastelli L, Rieger DK, Shenoy SG, Shir
Starling G, Spytek KA, Stone DJ, Tchernev VT,
, Zerhusen BD, Zhong M;
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2002US-0381495P.
2002US-0381495P.
2002US-0383534P.
2002US-03835744P.
2002US-0383829P.
2002US-0383829P.
2002US-0384024P.
2002US-0401783P.
2002US-0401753P.
2002US-00287971.
                                                                                                                                                                                                                                                                                                A,
                  (first entry)
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                                                                               protein;
                                                                                                                                                                                                                                                100.0%;
                                                                                   458
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                                                                                                                                                                                                                                                Score 117; DB 7;
Pred. No. 1.1e-09;
                                                                                 ₽
                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neuroprotective, antiparkinsonian,
                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcqueeney
Patturajan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shimkets
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15-MAY-2002;
17-MAY-2002;
28-MAY-2002;
28-MAY-2002;
28-MAY-2002;
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17-DEC-2001;
17-DEC-2001;
20-DEC-2001;
27-DEC-2001;
31-DEC-2001;
17-APR-2002;
15-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nootropic; neuroprotective; antiparkinsonian; anticonvulsant; osteopathic; antiarthritic; antiinflammatory; dermatological; antiasthmatic; antilipaemic; metabolic; diabetes; obesity; infectious; anorexia; cancer; cardiovascular; hypertension; attherosclerosis; neurodegenerative; Alzheimer's disease; Parkinson's; epilepsy; immune; osteoarthritis; haemopoietic; inflammatory skin; asthma; dyslipidaemia; neurogenesis; cell differentiation; proliferation; haemopoiesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NOVX; antidiabetic; anorectic; cardiant; hypotensive; antiarteriosclerotic; virucide; antibacterial; fungio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   healing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NOV15c
                                                                                                                                                                                                                                                                                                             2001US-0341477P
2001US-0341540P
2001US-0344297P
2001US-0344297P
2001US-0344297P
2002US-03732889
2002US-0380981P
2002US-0380981P
2002US-0381495P
2002US-0381495P
2002US-0381495P
2002US-0381495P
2002US-0381495P
2002US-0381495P
2002US-0381495P
2002US-03817971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001US-0336600P.
2001US-0338285P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001US-0341346P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   angiogenesis; gene therapy; chromosome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                fungicide; protozoacide;
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CURAGEN CORP.

Mezes PS, N Pena CEA, I Smithson G, Grosse WM, Lepley DM, Burgess Alsobrook JP, Ellerman Œ, JP, Alvarez E, Anderson DW, Baron M, Boldog FL;
, Casman SJ, Chapoval A, Dhanabal M, Edinger SR, Eisen;
, Ettenberg S, Gangolli EA, Gerlach VL, Gorman L;
Guo X, Hackett C, Ji W, Kekuda R, Khramtsov NV;
Li L, Macdougall JR, Malyankar UM, Mazur A, Mcqueeney
Miller CE, Millet I, Mishra VS, Padigaru M, Patturajan
Peyman JA, Rastelli L, Rieger DK, Shenoy SG, Shimkets R
Peyman JA, Rastelli L, Rieger DK, Tchernev VT, Twomlow
I, Zerhusen BD, Zhong M; Shimkets RA, ۶

2003-441555/41.

New isolated NOVX polypeptides and polynucleotides, useful for preventing, diagnosing or treating NOVX-associated disorders, e.g. osteoarthritis, obesity, atherosclerosis, cancer, Parkinson's disease, or infections

Claim SEQ ID ö 52; 447pp; English

The invention relates to a novel isolated NOVX polypeptide. The polypeptide of the invention demonstrates, antidabetic, anorect cardiant, hypotensive, antiarteriosclerotic, virucide, antibacte fungicide, protozoacide, nootropic, neuroprotective, antiparking anticonvulsant, osteopathic, antiarthritic, antiinflammatory, dermatological, antiasthmatic and antilipaemic activities. The polypeptides, nucleic acid molecules and antibodies may be useful treating or diagnosing diseases including metabolic disorders and antipathon of diagnosing diseases and dispersion of disorders and dispersion of dispersion disorders and dispersion disorders and dispersion d antiparkinsonian, c, anorectic, antibacterial, such for as

ភិទ្ធិសិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិត្ត សិក្សិ

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RESULT 12
AAW08349
ID AAW08
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XX AAW08
XX POORF
XX Recep
XX Recep
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XX Recep
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Best Local S
Matches 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Receptor antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiovascular diseases including hypertension and atherosclerosis, neurodegenerative disorders such as Alzheimer's disease, Parkinson's disease and epilepsy, immune disorders e.g. osteoarthritis, haemopoietic disorders, inflammatory skin disorders, asthma and dyslipidaemia. Furthermore, the nucleic acids and polypeptides may also be used to identify molecules that modulate or inhibit neurogenesis, cell differentiation and proliferation, haemopoietis, wound healing and angiogenesis, as well as in gene therapy. Finally, the nucleic acids may be used as hybridisation probes, in chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The current sequence is that of
                                                                                                                                    Recombinant immunogen corresp. to dimeric form of a receptor - generating antibodies able to act as receptor agonists, esp. of erythropoietin receptor for treating anaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
Chimeric.
                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EpoRFc fusion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW08349;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW08349 standard; protein; 488
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the human NOV protein of the invention.
                                                                                                                                                                                                                                                                                        WPI; 1997-051900/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9640231-A1
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                                                                                                                                                                                                                                                                                                                                                                                                  (SMIK ) SMITHKLINE BEECHAM CORP.
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                                                                                                                                                                                                                                                                                                                                             PR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QRVEILEGRTECVLSNLRGRTRY 23
                                                                                     Page 39-41;
                                                                                                                                                                                                                                                                                                                                             Erickson-Miller CL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; ilarity 100.0%; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          95US-00474673.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96WO-US009613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "erythropoietin receptor extracellular
251. .254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antibody; erythropoietin receptor; EpoR; immunogen;
nionein; promoter; IgGl; Fc; anaemia; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .488
                                                                                  83pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Factor Xa cleavage site"
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                                                                                     English.
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Pred. No. 1.1e-09;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IgG1
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sequence is that of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   458;
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                                                                                                                                                                                                  used
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A fusion protein (AAW08349) encoded by plasmid mtalsEpoRFc (AAT48800) comprises the human erythropoietin receptor (EpoR) extracellular domain

S

QRVEILEGRTECVLSNLRGRTRY 23

Matches

Query Match Best Local

Similarity

100.0%; llarity 100.0%; Conservative 0

0

Mismatches

Indels

<u>,,</u>

Gaps

0

Score 117; DB 3; Pred. No. 1.3e-09;

Length

503;

Sequence

503 AA;

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ARBSULT 13
ARAB13012
IID ARB13
XX ARB13
XX ARB13
XX O-tag
XX Site
XX Syntl
XX Tout
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           fused (via a Factor Xa cleavage sequence) to the Fc portion of human 19G1. It can be expressed e.g. in transfected Drosophila S2 cells upon induction with copper sulphate. The cells secrete EpoRFc as a dimeric molecule due to the affinity of the Fc moiety for itself. The dimeric receptor can be used as an immunogen to generate antibodies (monoclonal, polyclonal, chimeric, humanised) able to act as EcoR agonists for use in
                                    protein, and to a method for screening for a candidate compound which interacts with first protein. The screening method comprises contacting the candidate molecule with a labelled modified first protein and detecting the label to identify interaction of the labelled modified first protein and candidate compound. The first protein is modified to contain a peptide, represented by sequence ABB13005. The method is used to label proteins at specific sites. The present sequence represents a C tagged erythropoietin (EPO) receptor constructed in an example of the method of the invention
                                                                                                                                                                                                                                                                                           Screening for a candidate compound for use in bioassays comprises contacting the candidate molecule with a labelled modified protein detecting the label to identify interaction of the two molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    erythropoietin receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Site specific label; detection; interaction screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q-tagged
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                                                                                                                                                                                                                                                       Example 4; Page 26; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-JAN-2000; 2000WO-US001481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB13012 standard; protein; 503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of anaemia
                                                                                                                                                                                                                This invention relates to methods for the site specific modification
                                                                                                                                                                                                                                                                                         detecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200043492-A2
                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-499222/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         488 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            erythropoietin (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORVEILEGRIECVLSNLRGRIRY 216
                                                                                                                                                                                                                                                                                                                                                                                                              Powell DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0117327P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 117;
100.0%; Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                              Meek TD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                              Chen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                           modified protein two molecules.
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RESULT 14
AARO6512
ID AARO652
XX AARO652
AC AARO6
XX AARO
RESULT 15
AAR47518
ID AAR477
XX
AC AAR47
AC AAR47
DT 25-MA
DT 24-JU
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EYTH
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The sequence was deduced from DNA obtd. from a clone isolated from a commercially available human geenomic cDNA library in phage Lambda Fix (Strategene). The sequence encodes a type I trans-membrane protein with binding affinity for EPO. The gene and recombinant EPO receptor produced on expression of the DNA are used to develop reagents and systems to control and study erythropolesis. It is believed that the EPO receptor is dys-functional in individuals with Diamond Blackfan anaemia, and may be hyperactive in polycythemia vera. See also AARO6511 (murine EPO receptor). (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEMY )
Erythropoietin receptor; recombinant; murine; anaemia.
                                                                                                            25-MAR-2003
24-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Erythropoietin receptor and gene - used for developing reagents systems to control and study erythropoiesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
04-JAN-1991
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                                                      Human EPO receptor.
                                                                                                                                                                                             AAR47518;
                                                                                                                                                                                                                                                   AAR47518 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 508 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 2; 53pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dandrea A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-FEB-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-FEB-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Erythropoietin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EPO receptor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 al Similarity
23; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                          1 QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENETICS INST INC. WHITEHEAD INST BIOMEDICAL
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                                                                                                                                                                                                                                                                                                                                                                                       ORVEILEGRIECVLSNLRGRIRY 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wong G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Silarity 100.0%; Silarity 100.0%; Silarity 00.0%; Silarity 0;
                                                                                                          (revised)
(first entry)
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(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       89US-00306503.
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                                                                                                                                                                                                                                                protein; 508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 117; DB 2;
Pred. No. 1.3e-09;
; Mismatches 0;
                                                                                                                                                                                                                                                   B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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Search completed: May Job time : 55 secs
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                                                                                                                                                                         Matches
                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                              Mouse erythroleukaemia (MEL) cells were used to construct a cDNA library. The cDNA was used to transfect COS-1 cells and these were screened for radioiodinated erythropoietin (EPO) binding to isolate cDNA encoding the EPO receptor. This cDNA was used as a probe to screen a human genomic cDNA library to obtain DNA encoding the human EPO receptor. The cDNA may be used to study, treat or diagnose disorders in which the EPO receptor is dysfunctional. The EPO receptor may also be used to raise antibodied or for treating hypersensitivity to EPO or who have elevated levels of EPO. The pord, is gref, used for treating anaemias, primary proliferative polycythemia and secondary polycythemia. See also AAR47517. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                         Sequence 508 AA;
                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 9; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                      dysfunctional.
                                                                                                                                                                                                                                                                                                                                                                                                                   Recombinant DNA encoding erythropoietin receptor - used to develop prods. for study, treatment or diagnosis of disorders in which receptor is
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-025409/03.
N-PSDB; AAQ53995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        D'andrea A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEMY )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-FEB-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide
                                                          194 ORVEILEGRIECVLSNIRGRIRY 216
                                                                              1 QRVEILEGRTECVLSNLRGRTRY 23
                                                                                                                 l Similarity
23; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENETICS INST INC.
CHILDRENS MEDICAL CENT.
WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wong GG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             89US-00306503
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "putative transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "mature EPO receptor"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "signal peptide"
                6
                                                                                                               100.0%; Score 117; DB 2;
100.0%; Pred. No. 1.3e-09
tive 0; Mismatches 0
                 2004, 12:45:02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jones SS
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                                                                                                                                          Length 508;
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Minimum
Maximum
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Run on:
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BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     d. No. is the number of results predicted by chance to have a
re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           seq
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length: 2000000000
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Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              May 6, 2004, 12:42:22 ; Search time 23 Seconds (without alignments) 51.626 Million cell updates/sec
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Gapop 10.0 ,
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117
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1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/iaa/BACOMB.pep:*
6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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                                     Maximum Match 100%
Listing first 45 summaries
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                                 US-09-028-937-11
US-09-280-864A-12
US-09-022-291-12
US-09-339-838-7
US-09-339-838-7
US-08-950-293-5
US-08-950-293-5
US-09-134-000C-5287
US-09-132-210-1509
US-09-732-210-1600
US-09-732-210-1600
US-09-732-210-1601
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                                                                                                                                                                                                                                    Sequence 11, Appl
Sequence 12, Appl
Sequence 12, Appl
Sequence 5, Appli
Sequence 7, Appli
Sequence 7, Appli
Sequence 2, Appli
Sequence 4, Appli
Sequence 2, Appli
Sequence 2, Appli
Sequence 5287, Ap
Sequence 1600, Ap
Sequence 1500, Ap
Sequence 1203, A
Sequence 1203, A
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23930, Appli
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32222, A
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200, Ap
2067, A
2903, A
2903, A
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   , MOLECULE TYPE: protein US-09-028-937-11
                                                                                                                                                                                                                 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/701,382

FILING DATE: 22-AUG-1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/612,999

APPLICATION NUMBER: US 08/612,999

FILING DATE: 08-MAR-1996

ATTORNEY/AGENT INFORMATION:

NAME: Silva, Robin M.

REGISTRATION NUMBER: 38,304

REGISTRATION NUMBER: 38,304
                                                                                                                                TELEFAX: (415) 949-8711 INFORMATION FOR SEQ ID NO:
                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acids
TYPE: amino acid
STRANDENESS: unknown
                                                                                                                                                                  REFERENCE/DOCKET NUMBER: A-TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 781-1989
                                   TOPOLOGY:
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45	44	43	42	41	40	39	86	37	36	35	34	33	32	31	30	29	28
41	41	41	41	41	41	41	41	41	41	41	41	41	41	41	41	42	42
35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.0	35.9	35.9
1091	1018	1018	1018	1018	865	843	835	835	749	327	292	75	60	60	50	4545	4472
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US-08-986-485-5	US-08-040-741-6	US-08-714-901-6	US-08-408-420A-6	US-08-408-093-6	US-09-252-991A-20894	US-09-489-039A-7545	US-09-758-282B-243	US-09-758-282B-155	US-09-252-991A-17331	US-09-134-001C-3535	US-09-711-164-438	US-08-928-383B-13	US-08-754-477A-15	US-08-754-477A-14	US-09-621-976-6212	US-08-804-227C-14	US-08-804-227C-2
Seguence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence .	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
5, Appli	6, Appli	6, Appli	6, Appli	6, Appli	20894, A	7545, Ap	243, App	155, App	17331, A	3535, Ap	438, App	13, Appl	15, Appl	14, Appl	6212, Ap	14, Appl	719

ALIGNMENTS

US-09-028-937-11 Sequence 11, Applicati Patent No. 6333031 GENERAL INFORMATION: ZIP: 94111-4187 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatil FILING DATE: CLASSIFICATION: PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/788,820 FILING DATE: 23-JAN-1997 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA: NUMBER OF SEQUENCES: 4 CORRESPONDENCE ADDRESS: APPLICANT: Naranda, Tatjana TITLE OF INVENTION: Receptor Derived Peptides As Modulators TITLE OF INVENTION: Of Receptor Activity APPLICANT: ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert STREET: 4 Embarcadero Center, Suite 3400 CITY: San Francisco APPLICATION NUMBER: US/09/028,937 COUNTRY: ç Application US/09028937 USA Olsson, Lennart Version #1.30

unknown

A-63139-3/RFT/RMS

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RESULT 2
US-08-280-864A-12
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US-09-092-291-12
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Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Elliott, Steven G.
TITLE OF INVENTION: Antibodie
TITLE OF INVENTION: Erythorpo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Elliott, Steven G.
TITLE OF INVENTION: Antibodies Which Activat
TITLE OF INVENTION: Erythorpoietin Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Amgen Inc.
STREET: One Amgen Center
CITY: Thousand Oaks
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1840 CITY: Thousand Oaks
STATE: California
" S A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION: NAME: Winter, Robert B.
COMPUTER READABLE FORM
                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 30 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 424
                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23;
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               91320
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1840 Dehavilland Drive
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                            U.S.A.
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                                                                                                                                           Antibodies Which Activate 
Erythorpoietin Receptor
                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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Pred. No. 1.8e-12;
); Mismatches 0;
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US-09-339-838-7
Sequence 7, Application US/09339838
Patent No. 6361998
GENERAL INFORMATION:
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APPLICANT: Mueller, Susan G.
APPLICANT: Mueller, Susan G.
TITLE OF INVENTION: The Efficient Culture of
FILE REFERENCE: 6704-83
CURRENT APPLICATION NUMBER: US/09/339,838
CURRENT FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: CA 2,260,332
PRIOR APPLICATION NUMBER: CA 2,260,332
PRIOR FILING DATE: 1999-01-25
PRIOR APPLICATION NUMBER: CA 2,241,576
PRIOR FILING DATE: 1998-06-25
NUMBER OF SEQ ID NOS: 11
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                                                                        RESULT 5
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                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1 SEQ ID NO 5
                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                Query Match
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ATTORNEY/AGENT INFORMATION:
NAME: Winter, Robert B.
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                  LENGTH: 438
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 30 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA: APPLICATION NUMBER: US,
                                                                                                                                                                                                                   Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
COMPUTER: II
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                                                                                                                          194 QRVEILEGRTECVLSNLRGRTRY 216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 117; DB 4; 100.0%; Pred. No. 1.8e-12;
                                                                                                                                                                                                               100.0%; Score 117; DB 4; 100.0%; Pred. No. 4.3e-11;
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                                                                                                                                                                                                red. No. 4.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of Stem Cells for the Production of Hemoglok
                                                                                                                                                                                                                                    DB 4;
                                                                                                                                                                                                                                  Length 438;
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APPLICANT: Bell, David N.
APPLICANT: Mueller, Susan G.
APPLICANT: Matthews, Kathryn E.

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; LENGTH: 438
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-339-838-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-776-511-2
                                                                                                                               ; MOLECULE TYPE: protein US-08-776-511-2
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                                                         Matches
                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: The Efficient Culture of Stem Cells for the Production of Hemogld CURRENT APPLICATION NUMBER: US/09/339,838
CURRENT FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: CA 2,260,332
PRIOR FILING DATE: 1999-01-25
PRIOR FILING DATE: 1999-01-25
PRIOR APPLICATION NUMBER: CA 2,241,576
PRIOR APPLICATION STEM CA 2,241,576
PRIOR FILING DATE: 1998-06-25
NUMBER: OF SEQ ID NOS: 11
NUMBER: OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Young, Peter R.
APPLICANT: Erickson-Miller, Connie
TITLE OF INVENTION: Method for Obtaining Receptor Agonist
TITLE OF INVENTION: Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                             TELEFAX: 610-270-5090 INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 610-270-5015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                        NAME: Jervis, Herbert H.
REGISTRATION NUMBER: 31,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 709 Swedeland CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: SmithKline Beecham Corporation- Corporate ADDRESSEE: Patents
                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/776,511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
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QRVEILEGRTECVLSNLRGRTRY 23
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                                                       100.0%; Score 117; DB 3; ilarity 100.0%; Pred. No. 4.9e-11; Conservative 0; Mismatches 0;
                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                        424
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                                                                                             Length 488;
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                                                           Indels
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GENERAL INFORMATION:
APPLICANT: Lee, Jong Y.
TITLE OF INVENTION: PURIL
TITLE OF INVENTION: PROTI
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                           Sequence 4, Application US/09039609
Patent No. 6107473
                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/499,643
FILING DATE: 07-JUL-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/106,815
FILING DATE: 16-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Ellinger, Mark S.
REGISTRATION NUMBER: 34,812
REFERENCE/DOCKET NUMBER: 07004/0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612/335-5070
                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 612/288-9696 INFORMATION FOR SEQ ID NO:
                                                                           APPLICANT: ALBONE, EARL
APPLICANT: KIKLY, KRISTINE
TITLE OF INVENTION: A KRINGLE-RELATED CLONE,
TITLE OF INVENTION: HTHEZ47
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 508 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 60 South a
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                 STREET: P.O. Box your CITY: Valley Forge
                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               194 QRVEILEGRTECVLSNLRGRTRY 216
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                                         E: Ratner & Prestia
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COMPUTER: IBM compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSRO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/850,293
FILING DATE: 05-MAY-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Fish & Richardson P.C., P.A. 60 South Sixth Street, Suite 3300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROTEIN FRAGMENT AND ANTIBODIES DERIVED THEREFROM
                100.0%; Score 117; DB 2; 100.0%; Pred. No. 5.2e-11
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                                         DB 2;
                                       Length 508;
Indels
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Gaps
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CURRENT APPLICATION NUMBER: US/09/039,609

FILLING DATE: 16-MAR-1998

CLASSIFICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/060,623

FILLING DATE: 01-OCT-1997

ATTORNEY/AGENT INFORMATION:

NAME: Prestia, Paul F

REGISTRATION NUMBER: GP-70275

TELECOMMUNICATION INFORMATION:

TELEPHONE: 610-7407-0700
                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: ALBONE, EARL
APPLICANT: KIKLY, KRISTINE
TITLE OF INVENTION: A KRINGLE-RELATED CLONE,
TITLE OF INVENTION: HTHBZ47
             FILING DATE: 16-MAR-1998
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/66,623
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Prestia, Paul F
REGISTRATION NUMBER: 23,031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 321 amino acids
                                                                                                                                                COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/039,609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 4
CORRESSONDENCE ADDRESS
ADDRESSEE: Rather & Prestia
STREET: p.O. Box 980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                          STREET: F.C.
CITY: Valley Forge
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REFERENCE/DOCKET NUMBER: GP-70275
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ZIP: 19482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10;
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Similarity 50.0%;
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Pred. No. 4.1;
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CURRENT APPLICATION NUMBER: US/09/732,210
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: US 60/169,513
PRIOR FILING DATE: 1999-12-07
PRIOR APPLICATION NUMBER: US 60/169,340
PRIOR FILING DATE: 1999-12-07
NUMBER OF SEQ ID NOS: 1753
SEQ ID NO 1599
LENGTH: 67
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US-09-134-000C-5287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: ENTEROCOCCUS FABCALIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 032796-032
CURRENT APPLICATION NUMBER: US/09/134,000C
CURRENT APPLICATION NUMBER: US 60/055,778
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR FILING DATE: 1997-08-15
PRIOR FILING DATE: 1997-08-15
SOPTWARE: Patentin version 3.1
SEQ ID NOS: 6812
SOPTWARE: Patentin version 3.1
SEQ ID NO 5287
LENGTH: 64
TYPE: PRT
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Best Local Similarity 50.0
Matches 10; Conservative
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SEQUENCE CHARACTERISTICS:
LENGTH: 458 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                         Sequence 1599, Application US/09732210
Patent No. 6573361
GENERRAL INFORMATION:
APPLICANT: Bunkers, Greg J.
APPLICANT: Liang, Jihong
APPLICANT: Mittanck, Cindy A.
APPLICANT: Seale, Jeffrey W.
APPLICANT: Wu, Yonnie S.
TITLE OF INVENTION: Anti-fungal Proteins and Methods for Their Use
FILE REFERENCE: 38-21(15036)B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Enterococcus faecalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 610-407-0700
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nes 9; Conserv
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Query Match
Best Local Similarity
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Best Local Similarity
"hes 9; Conserve
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US-09-252-991A-32067
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                                                                                 ; ORGANISM: Pseudomonas aeruginosa US-09-252-991A-32067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-732-210-1600
                                                                                                                                                                                                                                                                                                                                                         Sequence 32067, Appli
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 1753
SEQ ID NO 1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1600, Application US/09732210 Patent No. 6573361
                                                                                                                                  SEQ ID NO 32067
LENGTH: 372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
           Matches
                                           Query Match
                                                                                                                                                                 APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: ARRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1990-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Wu, Yonnie S.

TITLE OF INVENTION: Anti-fungal Proteins and Methods for Their Use
FILE REFERENCE: 38-21 (15036) B

CURRENT APPLICATION NUMBER: US/09/732,210

CURRENT FILING DATE: 2000-12-07

PRIOR APPLICATION NUMBER: US 60/169,513

PRIOR PILING DATE: 1999-12-07

PRIOR PILING DATE: 1999-12-07

PRIOR PILING DATE: 1999-12-07

PRIOR PILING DATE: 1999-12-07

PRIOR PILING DATE: 1999-12-07
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APPLICANT: Liang, Jihong
APPLICANT: Mittanck, Cindy A.
APPLICANT: Seale, Jeffrey W.
APPLICANT: Wu, Yonnie S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Kluyveromyces lactis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 67
TYPE: PRT
ORGANISM: Kluyveromyces marxianus
Jes 12; Conservation
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                          Score 45; DB Pred. No. 22;
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Pred. No. 2;
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Pred. No.
         Mismatches
                                             DB 4;
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         6;
                                             Length 372;
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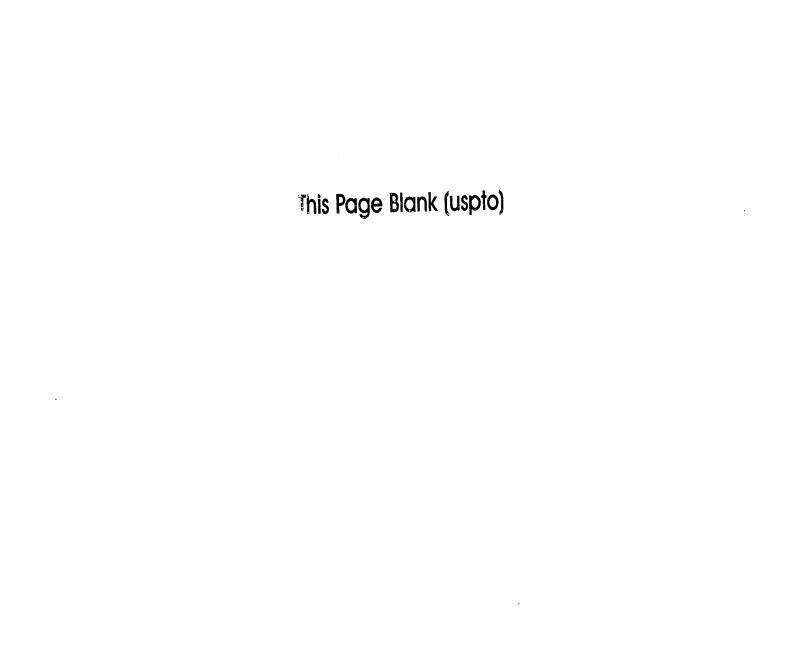
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 931
TYPE: PRT
ROANISM: Actinobacillus pleuropneumoniae
US-08-624-655A-2
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CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 12003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 12903, Application US/09489039A Patent No. 6610836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/08624655A Patent No. 6323005
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                                                                               Query Match
Best Local :
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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CURRENT APPLICATION NUMBER: US/08/624,655A
CURRENT FILING DATE: 1996-03-22
PRIOR APPLICATION NUMBER: 95 00592
PRIOR FILING DATE: 1995-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: DABAN, MONTSERRAT
APPLICANT: MEDRANO, ANDRES
APPLICANT: ESPUNA, ENRIC
APPLICANT: QUEROL, ENRIQUE
                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
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TYPE: PRT
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581
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                               6 LEGRIECVLSNLRGRIRY
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Pred. No. 45;
                                                                               Score 43; Pred. No. 1
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Search completed: May

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Job time : 23 secs



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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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| Cgn2 6/ptcdata/1/pubpaa/US07_PUBCOMB.pep:*
| Cgn2 6/ptcdata/1/pubpaa/PCT_NEW_PUB.pep:*
| Cgn2 6/ptcdata/1/pubpaa/US06_NEW_PUB.pep:*
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| Cgn2 6/ptcdata/1/pubpaa/US07_NEW_PUB.pep:*
| Cgn2 6/ptcdata/1/pubpaa/US07_NEW_PUB.pep:*
| Cgn2 6/ptcdata/1/pubpaa/PCTUS_PUBCOMB.pep:*
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Copyright (c) 1993 - 2004 Compugen Ltd.
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12 US-10-287-971-50
12 US-10-287-971-50
12 US-10-287-971-54
15 US-10-316-194-144
15 US-10-316-194-3
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          Sequence 11, Appl
Sequence 30, Appl
Sequence 48, Appl
Sequence 50, Appl
Sequence 52, Appl
Sequence 52, Appl
Sequence 54, Appl
Sequence 54, Appl
Sequence 142, App
Sequence 144, App
Sequence 144, App
Sequence 35, Appl
Sequence 35, Appl
Sequence 4, Appl
Sequence 14, Appl
Sequence 14, Appl
Sequence 11, Appl
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ALIGNMENTS

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CURRENT APPLICATION UNMEER: US/09/991,548
CURRENT FILING DATE: 2001-11-20
PRIOR FILING DATE: 2001-11-20
PRIOR FILING DATE: 1998-02-24
PRIOR FILING DATE: 1998-02-24
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: 08/701,382
PRIOR APPLICATION NUMBER: 08/612,999
PRIOR APPLICATION NUMBER: 08/612,999
PRIOR FILING DATE: 1996-08-22
PRIOR FILING DATE: 1996-03-08
NUMBER OF SEQ ID NOS: 44
SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 23
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APPLICANT: OLSSON, Lennart
APPLICANT: NARANDA, Tatjana
APPLICANT: NARANDA, Tatjana
TITLE OF INVENTION: RECEPTOR DERIVED PEPTIDES AS MODULATORS
TITLE OF INVENTION: OF RECEPTOR ACTIVITY
FILE REFERENCE: 213542000101
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                                                                                                                 Query Match
Best Local
                                                                                           Matches
                                                                                                                                                                                        FEATURE: OTHER INFORMATION: human erythropoietin receptor
                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                 Similarity
                        QRVEILEGRIECVLSNLRGRIRY 23
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                                                                                        Conservative
                                                                                                              100.0%; Score 117; DB 9; 100.0%; Pred. No. 1.8e-11;
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                                                                                           Mismatches
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RESULT 4
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Publication No. US20040023334A1
GENERAL INFORMATION:
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Publication No. US20030215444A1
GENERAL INFORMATION:
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30
LENGTH: 23
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Best Local S
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 12
LENGTH: 30
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TITLE OF INVENTION: Modified Transferrin Fusion Proteins
FILE REFERENCE: 54710-5001-US
CURRENT APPLICATION NUMBER: US/10/231,494
CURRENT FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: US 60/315,745
PRIOR PILING DATE: 2001-08-30
PRIOR PILING DATE: 2001-08-30
PRIOR PILING DATE: 2001-08-30
PRIOR PILING DATE: 2001-08-30
PRIOR PILING DATE: 2001-01-30
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PRIOR FILING DATE: 1994-07-26
PRIOR APPLICATION NUMBER: 09/640,090
PRIOR FILING DATE: 2000-08-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Elliott, Steven G
TITLE OF INVENTION: Antibodies which Activate an Erythropoietin Receptor
FILE REFERENCE: 06843-0030-04000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/092,671 PRIOR FILING DATE: 1998-06-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/364,276
CURRENT FILING DATE: 2003-06-17
                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: peptide
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Pred. No. 1.8e-11;
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                                                                                                                                                                                                       DB 15;
                                                                                                                                                                                                       Length
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US-10-287-971-50
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                                                                       SOFTWARE: CuraSeqList version 0.1 SEQ ID NO 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: CuraSeqList version 0.1 SEQ ID NO 48
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APPLICANT: Alsobrook, et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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PRIOR FILING DATE: 2002-08-26
NUMBER OF SEQ ID NOS: 397
                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/406,181 PRIOR FILING DATE: 2002-08-26
                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/348,283 PRIOR FILING DATE: 2001-11-09 PRIOR APPLICATION NUMBER: 60/393,262 PRIOR FILING DATE: 2002-07-02
                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/401,479
PRIOR FILING DATE: 2002-08-06
PRIOR APPLICATION NUMBER: 60/333,072
PRIOR FILING DATE: 2001-11-06
                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/997,425
PRIOR FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 10/035,568
PRIOR FILING DATE: 2001-10-22
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CURRENT FILING DATE: 2002-11-05
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PRIOR FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 10/035,568
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: 60/338,626
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CURRENT FILING DATE: 2002-11-05
                                                                                                                       NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Alsobrook, et al TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/333,072 PRIOR FILING DATE: 2001-11-06
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PRIOR APPLICATION NUMBER: 60/401,479
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ORGANISM: Homo sapiens
                           LENGTH: 458
TYPE: PRT
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US-09-016-159-5
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US-10-287-971-52
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PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: 60/338,626
PRIOR FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: 60/401,479
PRIOR APPLICATION NUMBER: 60/401,479
PRIOR FILING DATE: 2002-08-06
PRIOR APPLICATION NUMBER: 60/333,072
PRIOR APPLICATION NUMBER: 60/348,283
PRIOR FILING DATE: 2001-11-09
PRIOR FILING DATE: 2001-11-09
PRIOR FILING DATE: 2001-11-09
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SOFTWARE: CuraSeqList version 0.1
SEQ ID NO 52
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                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/09016159
Patent No. US20020031806A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/393,262
PRIOR FILING DATE: 2002-07-02
PRIOR APPLICATION NUMBER: 60/406,181
PRIOR FILING DATE: 2002-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/997,425
PRIOR FILING DATE: 2001-11-29
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CURRENT FILING DATE: 2002-11-05
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FILE REFERENCE: 21402-480A
ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                                                                                                                                                                                APPLICANT: Lee, Jong Y.
TITLE OF INVENTION: PURIFIED HUMAN ERYTHROPOIETIN RECEPTOR
TITLE OF INVENTION: PROTEIN FRAGMENT AND ANTIBODIES DERIVED THEREFROM
                                                                                                                                                             NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C., P.A.
STREET: 60 South Sixth Street, Suite 3300
                                                                                                                   STREET: 60 CONTROL OF TY: Minneapolis
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                                                                                                COUNTRY:
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                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/287,971
CURRENT FILING DATE: 2002-11-05
PRIOR APPLICATION UNMBER: 09/997,425
PRIOR FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 10/035,568
PRIOR FILING DATE: 2001-10-22
PRIOR PELING DATE: 2001-10-22
PRIOR PELING DATE: 2001-10-25
PRIOR PELICATION NUMBER: 60/338,626
PRIOR PILING DATE: 2001-11-05
PRIOR PILING DATE: 2001-11-05
PRIOR PILING DATE: 2002-08-06
PRIOR APPLICATION NUMBER: 60/333,072
PRIOR APPLICATION NUMBER: 60/3348,283
PRIOR PILING DATE: 2001-11-06
PRIOR APPLICATION NUMBER: 60/348,283
PRIOR PILING DATE: 2001-11-09
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                                            US-10-287-971-54
                                                                                                                              SOFTWARE: CuraSeqList
SEQ ID NO 54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 54, Applica Publication No. US20 GENERAL INFORMATION:
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Best Local Similarity
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Query Match
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PRIOR FILING DATE: 2002-08-26
NUMBER OF COLUMBER 2002-08-26
                                                                                                                                                                           NUMBER OF SEQ ID NOS: 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Alsobrook, et al TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES,
                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/393,262
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                                                               TYPE: PRT
ORGANISM: Homo
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APPLICATION NUMBER: US/09/016,159
FILING DATE: 30-JAN-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/734,097
FILING DATE: 21-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/460,525
                                                                                                            ENGTH:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 612/335-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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SOFTWARE: FastSEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Ellinger, Mark S. REGISTRATION NUMBER: 34
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GY: linear
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No. US20040067882A1
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                                                                 sapiens
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16-JUN-1997
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96.6%;
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Score 113;
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  DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEIC ACIDS ENCODING SAME, AND METHODS
Length 508;
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95.7%;

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CURRENT APPLICATION UNMBER: US/10/316,194
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 173
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 142
LENGTH: 109
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Query Match
Best Local Similarity
                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 173
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 144
                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 144, Application US/10316194
Publication No. US20030215914A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local (
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Matches
                                                                                                                                                                                                                                                                                                                                        APPLICANT: Houtzager, Erwin
APPLICANT: Vijn, Irma M.C.
APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
FILE REFERENCE: 2183-5610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -10-316-194-142
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APPLICANT: Vijn, Irma M.C.
APPLICANT: Sijmons, Peter C.
APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
FILE REFERENCE: 2183-5610US
                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/316,194
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
                                                                      FEATURE:
NAME/KEY: SITE
LOCATION: (1)..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: SITE
LOCATION: (1)..
                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: IMABIS027
                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: IMABIS020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                    ENGTH: 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               194 QRVKILEGRIECVLSNLRGRIRY 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 95.7
22; Conservative
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o. US20030215914A1
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                                                                      .. (109)
82.1%;
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Pred. No. 1.
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1; Mismatches 0
Score 96; DB 15;
Pred. No. 2.3e-07;
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                 Length 109;
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; Sequence 35, Application US/10316194
; Sequence No. US20030215914A1
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                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 43
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SEQ ID NO 35
LENGTH: 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 43, Application US/10316194
Publication No. US20030215914A1
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Query Match
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                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/316,194
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/316,194
CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 10/016,516
PRIOR FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 173
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Houtzager, Erwin
APPLICANT: Vijn, Irma M.C.
APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
FILE REFERENCE: 2183-5610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Vijn, Irma M.C.
APPLICANT: Sijmons, Peter C.
TITLE OF INVENTION: A structure for presenting desired peptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 2183-5610US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Houtzager, Erwin
                                                                                                                                                           TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: VAP amino acid
OTHER INFORMATION: sequence of iMab600
                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: VAP amino acid OTHER INFORMATION: sequence of iMab1100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence
                                                     LOCATION:
                                                                       NAME/KEY: SITE
                                                                                          FEATURE
                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: SITE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION:
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                                                     (1)..(109)
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87.0%;
 73.5%;
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 Score 86;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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 DB 15;
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 Length 109;
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SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 11
LENGTH: 1026
TYPE: PRT
ORGANISM: Homo sapiens
US-09-947-063-11
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RESULT 15
US-09-728-912-7
; Sequence 7, Application US/09728912
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US-09-947-063-4
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CURRENT APPLICATION NUMBER: US/09/947,063
CURRENT FILING DATE: 2001-09-05
PRIOR APPLICATION NUMBER: 60/229,990
PRIOR FILING DATE: 2000-09-05
PRIOR APPLICATION NUMBER: 60/229,988
PRIOR FILING DATE: 2000-09-05
PRIOR OF BEQ ID NOS: 30
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/09947063
Publication No. US20030059775A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 82. Matches 19; Conservative
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Publication No. US20030059775A1
GENERAL INFORMATION:
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                                                                                                                                                                     Matches
                                                                                                                                                                                  Query Match 44.4%;
Best Local Similarity 71.4%;
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                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Padigaru et al.
TITLE OF INVENTION: No. US20030059775A1el Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-112
CURRENT APPLICATION NUMBER: US/09/947,063
CURRENT FILING DATE: 2001-09-05
PRIOR APPLICATION NUMBER: 60/229,990
PRIOR APPLICATION NUMBER: 60/229,988
PRIOR APPLICATION NUMBER: 60/229,988
PRIOR PILING DATE: 2000-09-05
PRIOR FILING DATE: 2000-09-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Padigaru et al.
TITLE OF INVENTION: No. US20030059775Alel Proteins and Nucleic Acids Encoding Same
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TYPE: PRT
ORGANISM: Homo sapiens
                                                                                             292
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                                                                                                                                                                     10; Conservative
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                                                                                                                            7 EGRTECVLSNLRGR 20
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                                                                                             EGFYECIASNLRGR 305
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71.48;
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Pred. No. 27;
                                                                                                                                                                   Score 52; DB 10;
Pred. No. 27;
1; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 8.6e-06; 0; Mismatches 4.
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Search completed: May Job time : 43 secs
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CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 60/169,252
PRIOR FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. US20010036643A1
GENERAL INFORMATION:
                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Holloway, James L.
TITLE OF INVENTION: Tumor Necrosis Factor-Stimulated Gene
TITLE OF INVENTION: and Protein
FILE REFERENCE: 99-94US
                                                                                                                                                                                                                       LENGTH: 49
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                       10;
                                                                                     3 VEILEGRIECVLSNLRGRIR 22
                                                                   7 VELLDGYTHRVLARFHGRSR 26
                                                                                                                                                                                                                                                                                         FastSEQ for Windows Version 3.0
                                                                                                                                       Conservative
                6, 2004, 12:48:22
                                                                                                                                                     41.9%;
                                                                                                                                       4; Mismatches
                                                                                                                                                     Score 49; I
Pred. No. 2.
                                                                                                                                                                       DB 9;
                                                                                                                                       6; Indels
                                                                                                                                                                       Length 49;
                                                                                                                                       0
                                                                                                                                       Gaps
```

0,

